

(19) World Intellectual Property Organization
International Bureau



(43) International Publication Date
3 April 2003 (03.04.2003)

PCT

(10) International Publication Number
WO 03/027323 A1

(51) International Patent Classification⁷: **C12Q 1/68**

(21) International Application Number: PCT/KR01/01562

(22) International Filing Date:
18 September 2001 (18.09.2001)

(25) Filing Language: Korean

(26) Publication Language: English

(30) Priority Data:
10-2001-0056827
14 September 2001 (14.09.2001) KR

(71) Applicant (*for all designated States except US*):
BIOMEDLAB CO., LTD. [KR/KR]; 4F., Dongsung
Bldg., 1-49 Dongsung-dong, Jongro-ku, Seoul 110-510
(KR).

(72) Inventors; and

(75) Inventors/Applicants (*for US only*): **YOON, Sung-Wook**
[KR/KR]; RM. 501, Munseong Bldg., 547-3, Dogok-dong,

Kangnam-ku, Seoul 135-270 (KR). **PARK, Tae-Shin**
[KR/KR]; 81-4, Yeonhui 2-dong, Seodaemun-ku, Seoul
120-113 (KR). **KIM, Joeng-Mi** [KR/KR]; 7-208, Mis-
ung Apt., Bulgwang 1-dong, Eunpyong-ku, Seoul
122-041 (KR). **PARK, Mi-Sun** [KR/KR]; 316, Jure-dong,
Sasang-ku, Busan 617-010 (KR).

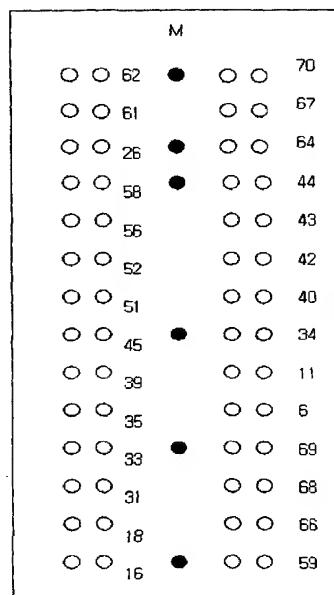
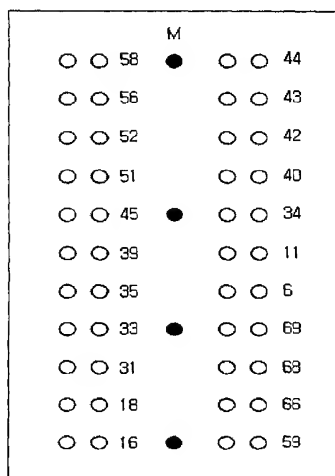
(74) Agent: **YOU ME PATENT & LAW FIRM**; Teheran
Bldg., 825-33, Teheran Bldg., 825-33, Yoksam-dong,
Kangnam-ku, Seoul 135-080 (KR).

(81) Designated States (*national*): AE, AG, AL, AM, AT, AU,
AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU,
CZ, DE, DK, DM, DZ, EC, EE, ES, FI, GB, GD, GE, GH,
GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KZ, LC, LK,
LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX,
MZ, NO, NZ, PH, PL, PT, RO, RU, SD, SE, SG, SI, SK,
SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA,
ZW.

(84) Designated States (*regional*): ARIPO patent (GH, GM,
KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian
patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European
patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE,

[Continued on next page]

(54) Title: GENOTYPING KIT FOR DIAGNOSIS OF HUMAN PAPILLOMA VIRUS INFECTION



(57) Abstract: The present invention relates to a genotyping kit for diagnosis of detecting the human papillomavirus (HPV) infection, probes for genotyping the HPV, and DNA chips including the probes. Also, the present invention relates to a method for diagnosis of HPV infection.

WO 03/027323 A1



IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, ML, MR, NE, SN, TD, TG).

— *with amended claims and statement*

Published:

— *with international search report*

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

GENOTYPING KIT FOR DIAGNOSIS OF HUMAN PAPILLOMA VIRUS INFECTION

BACKGROUND OF THE INVENTION

5 (a) Field of the Invention

The present invention relates to a genotyping kit and method for diagnosis of human papillomavirus (HPV) infection, probes for genotyping the HPV, and DNA chips including the probes. More specifically, the present invention relates to a genotyping kit for detecting human papillomaviruses from
10 clinical samples of infected patients using a DNA chip, a process for preparing the said DNA chip, and a method for diagnosis of HPV infection using the genotyping kit.

(b) Description of the Related Art

15 Uterine cancer includes cervical cancer, endometrial cancer, uterine sarcoma and the like. For cervical cancer, approximately 450,000 new cases occur worldwide each year, approximately 6,000 of these in Korea. Since the occurrence of cervical cancer (including cervical intraepithelial neoplasia) accounts for 22.1% of total cancer cases in Korean women, the highest
20 incidence with the second highest death rate, the prevention, diagnosis and treatment of cervical cancer are regarded as the most important issues in women's health.

Cervical cancer progresses through a precancerous stage, cervical intraepithelial neoplasia (CIN), known to be mainly caused by human

papillomavirus (HPV) infection. Especially, infection by particular types of HPV raises the possibility of developing invasive disease. Over 70 genotypes of HPV have been identified since the recognition of HPV as the main etiological factor for cervical cancer, and through research, certain HPV
5 genotypes have been selectively found in lesions of specific locations or progression stages, which has furthered recognition of the biological diversity of HPV infection. Among the HPV genotypes detected in the anogenital area, over 10 genotypes have been classified as being part of the high-risk group that are associated with an elevated risk for developing cervical cancer.
10 Based on these findings, characterization of the biological differences of HPV infection is considered to be of significant importance to the diagnosis and prevention of cervical cancer.

The test that has been most commonly used for the diagnosis of cervical cancer at its early stage is the Pap smear, which is a cytological test
15 performed as follows: old cells removed from the outermost layer of cells from the surface of the cervix are stained and examined for histopathological characteristics of HPV infection including koilocytosis, perinuclear halo formation in the epithelial cells. However, due to the low diagnostic efficiency (1 - 15%) of Pap tests together with other limitations, additional methods such
20 as colposcopy are necessary for more dependable diagnoses. Colposcopic screening can detect HPV infection at up to 70%, but it has disadvantages including high cost of the equipment, the need for skilled interpreters, and inability of determining HPV genotypes to distinguish between high-risk and low-risk infection. Therefore, efforts have been continuously made to develop

techniques for the detection of HPV and identification of HPV genotypes, to supplement conventional screening methods for cervical cancer and its precursors such as the Pap test.

The methods for detection of HPV and identification of HPV genotypes
5 can be classified into two groups, i.e., direct detection of HPV DNA and
detection of amplified HPV DNA. The methods for direct detection of HPV
DNA include liquid hybridization (Hybrid Capture kit by Digene Diagnostics,
Silver Spring, MD, USA, www.digene.com), Southern blot and dot blot with
HPV type-specific probes, filter in situ hybridization (FISH) and the like; and
10 methods for the detection of amplified DNA include type-specific PCR
(polymerase chain reaction) and general-primer PCR. In particular, genotype
analyses of amplified HPV DNA by general primer sets are commonly
performed by employing dot blot hybridization, microtitre plate hybridization, or
a line probe assay. Among these methods, liquid hybridization by Hybrid
15 Capture and line probe assay following a general-primer PCR have been
considered most suitable for diagnostic purposes. The line probe assay can
detect about 20 different HPV genotypes by immobilizing oligonucleotide
probes on a nitrocellulose membrane, however it lacks reliability due to low
sensitivity and difficulties in data interpretation. Commercialized Hybrid
20 Capture kits can detect HPV DNA in clinical samples without PCR amplification,
and they can distinguish between high-risk and low-risk HPV groups. But the
fact that Hybrid Capture kits cannot identify the genotypes of infecting HPV
limits accurate risk determination since the risk factors amongst high-risk HPVs
are not the same, in other words, intermediate-risk types are included in the

high-risk group. Moreover, the use of an RNA probe may present low stability of the kit, and the possibility of contamination cannot be excluded.

Under these circumstances, there have been strong reasons for exploring and developing a simple and accurate method and a genotyping kit
5 for detection of HPV infection and identification of the genotype of infecting HPV with high specificity and sensitivity, as well as probes for genotyping the HPV, and DNA chips including the probes. In addition, the genotyping kit, the probes and the DNA chips must have specificity and sensitivity high enough to detect various kinds of HPV so that they can diagnose the HPV infection and
10 identify its genotype with accuracy.

SUMMARY OF THE INVENTION

It is one object of the present invention to provide novel probes having nucleotide sequences complementary to DNA of HPV in order to detect HPV
15 infection and identify the genotype of HPV with high specificity and sensitivity.

It is another object of the invention to provide a DNA chip for detection of HPV infection and identification of the HPV genotype, wherein the DNA chip comprises probes having nucleotide sequences complementary to DNA of HPV.

It is yet another object to provide an HPV genotyping kit comprising a
20 novel DNA chip for detection of HPV infection and identification of the HPV genotype.

It is still another object to provide a method for diagnosis of HPV infection and identification of the HPV genotype using a novel DNA chip for detection of HPV infection and identification of the HPV genotype.

It is a further object to provide a method for amplifying sample DNA by a novel PCR in the diagnosis of HPV infection and the identification of the HPV genotype using a novel DNA chip for detection of HPV infection and identification of the HPV genotype.

5

BRIEF DESCRIPTION OF THE DRAWINGS

Figures 1a and 1b are schematic representations of the types and positions of the probes on the DNA chip according to one embodiment of the present invention.

Figure 2a is a photograph showing the result of HPV 16 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2b is a photograph showing the result of HPV 18 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2c is a photograph showing the result of HPV 39 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2d is a photograph showing the result of HPV 58 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2e is a photograph showing the result of HPV 68 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2f is a photograph showing the result of HPV 69 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2g is a photograph showing the result of HPV 6 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2h is a photograph showing the result of HPV 11 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2i is a photograph showing the result of HPV 43 DNA analysis using the DNA chip shown in Figure 1a.

Figure 2j is a photograph showing the result of HPV 44 DNA analysis using the DNA chip shown in Figure 1a.

5 Figure 2k is a photograph showing the result of HPV 61 DNA analysis using the DNA chip shown in Figure 1b.

Figure 2l is a photograph showing the result of HPV 67 DNA analysis using the DNA chip shown in Figure 1b.

10 Figure 3a is a photograph showing the result of analyzing a sample infected with HPV 16 using the DNA chip of the present invention.

Figure 3b is a photograph showing the result of analyzing a sample infected with HPV 31 using the DNA chip of the present invention.

Figure 3c is a photograph showing the result of analyzing a sample infected with HPV 51 using the DNA chip of the present invention.

15 Figure 3d is a photograph showing the result of analyzing a sample infected with HPV 52 using the DNA chip of the present invention.

Figure 3e is a photograph showing the result of analyzing a sample infected with HPV 56 using the DNA chip of the present invention.

20 Figure 3f is a photograph showing the result of analyzing a sample double-infected with HPV 16 and HPV 68 using the DNA chip of the present invention.

Figure 3g is a photograph showing the result of analyzing a sample double-infected with HPV 16 and HPV 58 using the DNA chip of the present invention.

Figure 3h is a photograph showing the result of analyzing a sample double-infected with HPV 33 and HPV 56 using the DNA chip of the present invention.

5 Figure 3i is a photograph showing the result of analyzing a sample triple-infected with HPV 16, HPV 52 and HPV 59 using the DNA chip of the present invention.

Figure 3j is a photograph showing the result of analyzing a sample triple-infected with HPV 16, HPV 35 and HPV 59 using the DNA chip of the present invention.

10 Figure 4 is a photograph showing the result of DNA amplification by two-step PCR according to one preferred embodiment of the present invention.

Figure 5 is a photograph showing the result of DNA amplification by multiplex PCR according to one preferred embodiment of the present invention.

15 Figure 6 is a schematic representation of the types and positions of the probes of the present invention and the probes of prior art.

Figure 7a is a photograph showing the result of HPV 16 DNA analysis using the DNA chip shown in Figure 6.

Figure 7b is a photograph showing the result of HPV 18 DNA analysis using the DNA chip shown in Figure 6.

20 Figure 7c is a photograph showing the result of HPV 56 DNA analysis using the DNA chip shown in Figure 6.

Figure 7d is a photograph showing the result of HPV 43 DNA analysis using the DNA chip shown in Figure 6.

Figure 7e is a photograph showing the result of HPV 68 DNA analysis

using the DNA chip shown in Figure 6.

Figure 8 is a photograph showing the result of HPV 66 DNA analysis using the HPV 66 probes of the present invention and the HPV 66 probes of the prior art.

5

DETAILED DESCRIPTION OF THE PREFERRED EMBODIMENTS

The genotyping kit of the invention for diagnosis of human papillomavirus (HPV) infection comprises: a DNA chip with probes that have nucleotide sequences complementary to DNA of HPV; primers for amplifying DNA obtained from clinical samples by PCR; and means for labeling amplified DNA hybridized with the probes of the said DNA chip.

The present inventors have tried to detect HPV infection and identify the types of HPV by way of genotyping DNA from clinical samples and prepared probes that have nucleotide sequences complementary to the DNA of HPV, and a DNA chip including the said probes. Further, we prepared an HPV genotyping kit comprising a DNA chip including probes that have nucleotide sequences complementary to the DNA of HPV, primers for amplifying DNA obtained from clinical samples by PCR, and means for labeling amplified DNA hybridized to the probes of the said DNA chip, and we successfully detected HPV infection and identified genotypes of infecting HPV with the aid of the genotyping kit in a simple and accurate manner.

The present invention provides nucleotide sequences of HPV DNA, preferably nucleotide sequences complementary to the L1 region of HPV DNA

as probes. The probes of the present invention comprise nucleotide sequences set forth in SEQ ID NO: 1-30 (Tables 1a and 1b). In Table 1, the underlined regions on SEQ ID NO: 25 are additional sequences to provide the conformation suitable for the function of probes.

5

[Table 1a]

SEQ ID NO.	HPV Type	Sequence
Probes for newly added HPV Types		
1	13	5'-ACATCATCTCTTTCAGACACATATAAGGCC-3'
2	26	5'-AGTACATTATCTGCAGCATCTGCATCCACT-3'
3	30	5'-TTATCCACATATAATTCAAGCCAAATTAAA-3'
4	43	5'-CCTCTACTGACCCTACTGTGCCAGTACAT-3'
5	53	5'-TCTACATATAATTCAAAGCAAATTAAACAGTA-3'
6	54	5'-TACAGCATCCACGCAGGATAGCTTTAATAAT-3'
7	54	5'-CATCCACGCAGGATAGCTTTAATAATTCTG-3'
8	55	5'-TGTGCTGCTACAACCTCAGTCTCCATCTACAACA-3'
9	57	5'-GAAACTAATTATAAAGCCTCCAATTATAAGGAA-3'
10	61	5'-CTGTATCTGAATATAAAGCCACAAGCTTTAG-3'
11	61	5'-CCTGTATCTGAATATAAAGCCACAAGCTTT-3'
12	62	5'-CCTCCACTGCTGCAGCAGAATACACGGCTA-3'
13	64	5'-TACAAATCCACCATATGCAAACACTAATTTTAA-3'
14	67	5'-TATGTTCTGAGGAAAAATCAGAGGCTACAT-3'
15	68	5'-TTTGTCTACTACTACTGAATCAGCTGTACCAA-3'
16	69	5'-AATCTGCATCTGCCACTTTTAAACCATCAGATT-3'
17	70	5'-AACGGCCATACCTGCTGTATATAGCCCTAC-3'
18	70	5'-ACCGAAACGGCCATACCTGCTGTATATAGC-3'
19	74	5'-CGCCTTCTGCTACATATAATAGTTCAGACT-3'
20	JC9710	5'-AAACACCCTCTGACACATACAAGGCTTCCA-3'
21	16	5'-GTGCTGCCATATCTACTTCAGAAACTACAT-3'
22	16	5'-TATGTGCTGCCATATCTACTTCAGAAACTACATA-3'
23	34	5'-GGTACACAATCCACAAGTACAACCTGCACCA-3'
24	35	5'-TTCTGCTGTGTCTTCTAGTGACAGTACATA-3'

25	35	5'-GCACGGTCTGTGTGTTCTGCTGTGTCTTCTACGTGC-3'
26	40	5'-CTTATGTGCTGCCACACAGTCCCCACACC-3'
27	56	5'-TATTAGTACTGCTACAGAACAGTTAAGTAA-3'
28	58	5'-CACTGAAGTAACTAAGGAAGGTACATATAA-3'
29	59	5'-TCTACTACTTCTTCTATTCTTAATGTATAC-3'
30	66	5'-CTAAAAGCACATTAATAATATGATGCCC-3'

[Table 1b]

31	6	5'-ATCCGTAACATCATCTTCCACATACACCAA-3'
32	11	5'-ATCTGTGTCTAAATCTGCTACATACACTAA-3'
33	18	5'-TGCTTCTACACAGTCTCCTGTACCTGGGCA-3'
34	31	5'-TGTTTGTGCTGCAATTGCAAACAGTGATAC-3'
35	33	5'-TTTATGCACACAAGTAACTAGTGACAGTAC-3'
36	39	5'-TCTACCTCTATAGAGTCTTCCATACCTTCT-3'
37	42	5'-CTGCAACATCTGGTGATACATATACAGCTG-3'
38	44	5'-GCCACTACACAGTCCCCTCCGTCTACATAT-3'
39	45	5'-ACACAAAATCCTGTGCCAAGTACATATGAC-3'
40	51	5'-AGCACTGCCACTGCTGCGGTTTCCCCAACA-3'
41	52	5'-TGCTGAGGTTAAAAAGGAAAGCACATATAA-3'
42	β -globin	5' -TGCACCTGACTCCTGAGGAGAAGTCTGCCG -3'
47	16	5'-GTCATTATGTGCTGCCATATCTACTTCAGA-3'
48	34	5'-TACACAATCCACAAGTACAAATGCACCATA-3'
49	35	5'-GTCTGTGTGTTCTGCTGTGTCTTCTAGTGA-3'
50	40	5'-GCTGCCACACAGTCCCCACACCAACCCCA-3'
51	56	5'-GTAAGTCTACAGAACAGTTAAGTAAATATG-3'
52	58	5'-ATTATGCACTGAAGTAACTAAGGAAGGTAC-3'
53	59	5'-CTGTGTGTGCTTCTACTACTGCTTCTATTCT-3'
54	66	5'-CTATTAATGCAGCTAAAAGCACATTAATA-3'

The probes of the present invention enable highly specific and sensitive detection of HPV infection and identification of the HPV genotype,

and have higher specificity and sensitivity than the probes for HPV genotyping described in PCT/KR/01213 (filed on October 26, 2000). Among the probes of the present invention, the probes set forth in SEQ ID NO: 1-20, HPV 13, 26, 30, 43, 53, 54, 55, 57, 61, 62, 64, 67, 68, 69, 70, 74, and JC9710 are not
5 included in the genotyping kit described in the above application, and therefore the probes according to the present invention enable it to be an implement that can detect more various types of HPV, contributing to its accuracy. The probes for HPV genotyping described in PCT/KR/01213 further comprise additional probes for HPV 16, 34, 35, 40, 56, 58, 59, and 66, along with
10 nucleotide sequences set forth in SEQ ID NO: 31-41.

The DNA chip of the present invention may comprise one or more HPV probes selected from nucleotide sequences set forth in SEQ ID NO: 1-30. In addition, it may further comprise a β -globin sequence as a marker, and preferably the nucleotide sequence set forth in SEQ ID NO: 42. In one
15 embodiment of the invention, the DNA chip comprises DNA chips including all probe sequence sets that are easily combined by those skilled in the art. It includes the following.

Embodiment 1

A DNA chip comprising one or more HPV probes selected from the
20 group consisting of oligonucleotides having nucleotide sequences set forth in SEQ ID NO: 1 to 30 and oligonucleotides having nucleotide sequences complementary to the said oligonucleotides.

Embodiment 2

A DNA chip comprising:

(i) one or more HPV probes selected from the group consisting of oligonucleotides having nucleotide sequences set forth in SEQ ID NO: 1 to 30 and oligonucleotides having nucleotide sequences complementary to the said oligonucleotides; and

5 (ii) one or more HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

Embodiment 3

A DNA chip comprising:

10 (i) one or more HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 1 to 20 and nucleotide sequences complementary to the said nucleotide sequences;

(ii) 8 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide sequences complementary to the said nucleotide sequences; and

15 (iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

Embodiment 4

20 A DNA chip comprising:

(i) 9 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 4 to 10;

(ii) 8 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide

sequences complementary to the said nucleotide sequences; and

(iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

5

Embodiment 5

A DNA chip comprising:

(i) 3 HPV probes having nucleotide sequences set forth in SEQ ID NO: 4, 15, and 16;

10

(ii) 8 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide sequences complementary to the said nucleotide sequences; and

(iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

15

In one embodiment of the invention, the DNA chip may further comprise position markers to locate probes. The position markers include, but are not limited to, β -globin, actin, and GAPDH (glyceraldehyde-3-phosphate dehydrogenase) genes, and the like. The preferable position marker is β -globin, whose sequence is set forth in SEQ ID NO: 42.

20

According to the present invention, probes that have nucleotides complementary to DNA of HPV are provided. The probes of the invention are characterized to specifically hybridize with DNA of HPV.

The process for preparing a DNA chip contained in the said HPV genotyping kit comprises the steps of: preparing 5' terminal amine-linked DNA

probes which have nucleotide sequences complementary to DNA of HPV; affixing the DNA probes thus prepared to an aldehyde-derivatized solid surface; and reducing excessive aldehydes not reacted with amine. The process for preparing a DNA chip of the invention is described in more detail by
5 the following steps.

Step 1: Preparation of probes

5' terminal amine-linked DNA probes that have nucleotide sequences complementary to the DNA of HPV are prepared. The nucleotide sequences of the probes are designed and synthesized to have nucleotide sequences
10 complementary to the DNA of HPV, preferably the L1 region of HPV DNA, and the probes are prepared by linking an amine group at the 5' terminal of the nucleotide sequences which enables the probes to bind to an aldehyde-derivatized solid surface.

Step 2: Affixture of probes

15 DNA probes prepared in Step 1 are affixed to an aldehyde-derivatized surface of a solid support, preferably glass. The probes are affixed to the surface of the solid support via a Schiff's base reaction between an aldehyde group on the surface of the solid support and an amine group at the 5' terminal of the probe under the condition of a 30 to 40°C temperature and 70 to 100%
20 humidity, while controlling the concentration of probes in a range of preferably 100 to 300 pmol/ μ l, and more preferably 200 pmol/ μ l. Also, β -globin which functions as a control for position markers and the hybridization reaction are affixed to the surface of the solid support.

Step 3: Preparation of DNA chip

Excessive aldehydes not reacted with the amine on the solid surface are reduced by employing a reducing agent of NaBH_4 , to finally prepare a DNA chip.

5 The present invention also provides an HPV genotyping kit comprising the said DNA chip and a method for diagnosis of HPV infection using the said genotyping kit. The present invention relates to an HPV genotyping kit comprising:

- (i) a DNA chip according to the invention;
- (ii) primers for amplifying DNA obtained from clinical samples by
10 PCR; and,
- (iii) means for labeling amplified DNA hybridized with the said DNA chip.

The present invention also relates to a method for diagnosis of HPV infection comprising the steps of:

- 15 (i) amplifying DNA obtained from clinical samples by PCR with primers of the HPV genotyping kit, for example primers set forth in SEQ ID NO: 43 and 44;
- (ii) applying the amplified DNA to the DNA chip of the HPV genotyping kit to hybridize the amplified DNA with probes of the DNA chip; and
- 20 (iii) detecting DNA bound on the surface of the DNA chip after labeling hybridized DNA.

In one embodiment according to the invention, amplification of sample DNA may be carried out via a usual PCR, preferably a two-step PCR or a multiplex PCR that amplifies using more than 2 primers at the same time, and

more preferably a multiplex two-step PCR that amplifies in two steps using more than 2 primers.

The development of the best condition for the HPV DNA sequence amplification methodology resulted in the two-step PCR or the multiplex PCR.

5 PCR consists of pre-denaturation, denaturation, primer annealing, and chain extension.

 While conventional PCR is performed by repeating one step under the same conditions, two-step PCR is performed with another amplification step, that is, two-step PCR is performed by repeating one or more steps among
10 denaturation, primer annealing, or chain extension under different conditions. According to the present invention, the first amplification step is carried out for a shorter time than that of the conventional chain extension step, and the second amplification step is carried out for an even shorter time than that of the first amplification step. Therefore, it prevents obtaining lengthy products when
15 the extension reaction is carried out for a longer time (e.g. 1 to 2 minutes), and thus increases the specificity of probes, which is preferable. More preferably, a chain extension of the PCR amplification may be carried out for 20 to 30 seconds in the first step, and for 10 to 15 seconds in the second step.

 In the present invention, while it is possible to amplify the HPV primer
20 sample, the β -globin sample, and a marker separately, one-step amplification using HPV primers set forth in SEQ ID NO: 43-44 and β -globin primers set forth in SEQ ID NO: 45-46 can be performed together. Otherwise, two-step PCR as mentioned above can also be done.

 PCR amplification may be carried out using an HPV primer sample

and a β -globin sample, respectively, or it may be carried out in one step or two steps using HPV primers set forth in SEQ ID NO: 43-44 and β -globin primers set forth in SEQ ID NO: 45-46, concomitantly.

In one preferred embodiment of the invention, PCR was performed in a PCR thermocycler containing the said primers and β -globin, dATP, dCTP, dGTP, dTTP, labeling materials, and *Taq* polymerase, and optionally BSA and DMSO, with 5 cycles of denaturation for 5 min at 94 °C, denaturation for 1 min at 94 °C, primer annealing for 2 min at 45~50 °C, and extension for 20~30 sec at 72 °C, followed by 35 cycles of denaturation for 1 min at 94 °C, primer annealing for 2 min at 45~50 °C and extension for 10~15 sec at 72 °C, and then further extension for 2 min at 72 °C, resulting in obtaining labeled amplified DNA samples.

The method for diagnosis of HPV infection using HPV genotyping kit of the invention is further illustrated by the following steps.

Step 1: Amplification of sample DNA

DNA obtained from clinical samples is amplified using the primers of the HPV genotyping kit, where polymerase chain reaction (PCR) employing biotin-16-dUTP is carried out to give biotin-containing amplified DNA. 4 primers for amplification are set forth in Table 2:

[Table 2]

SEQ ID NO.	HPV Type	Sequence	Base Number
43	Gp5+	5'-TTTKTTACHGTKGTDGATACYAC-3'	23
44	Gp6+	5'-GAAAHATAAAYTGYYAADTCATAYTC-3'	25
45	BG 1	5'-ATACAAGTCAGGGCAGAG-3'	18
46	BG 2	5'-CTTAAACCTGTCTTGTAACC-3'	20

In Table 1, R(A, g): Y(C, T): M(A, C): K(g, T): S(g, C): W(A, T), V(A, C, g): H(A, T, C): B(g, T, C): D(g, A, T): N(A, g, V, T), which is well-known to those skilled in the art.

Primers for DNA amplifying composed of nucleotide sequences set forth in SEQ ID NO: 43 and 44 are provided, as well as novel β -globin primers composed of nucleotide sequences set forth in SEQ ID NO: 45 and 46.

Step 2: Hybridization

Amplified DNA thus obtained is applied to the DNA chip of the HPV genotyping kit and hybridized with the probes of the DNA chip.

Step 3: Detection

The amplified sample DNA hybridized with the probes are labeled with means for labeling and detected with a confocal laser scanner.

Means for labeling comprise, but are not limited to, Cy5, biotin-binding compound, Cy3, EDANS (5-(2'-aminoethyl)amino-1-naphthalene sulfuric acid), tetramethylrhodamine (TMR), tetramethylrhodamine isocyanate (TMRITC), x-rhodamine or Texas red. Preferably, Cy5 and biotin-binding compound can be used. Labeling materials are detected by a confocal laser scanner.

The HPV genotyping kit of the invention is an implement that can

detect HPV infection in a simple and accurate manner, as well as identify the types of infecting HPV, and therefore it may contribute to early diagnosis, prevention, and treatment of cervical cancer.

The present invention is further shown in the following examples, which should not be taken to limit the scope of the invention. Particularly, though a DNA chip with 22 or 28 probes was prepared in the Examples described below, it is to be understood that the present invention is not limited by types and numbers of probes, but DNA chips using nucleotide sequences derived from HPV DNA and any variety of detection kits using the said DNA chips are intended to be included within the scope of the invention.

Example 1: Preparation of Probes

Prevalent HPV types including 15 high-risk types (HPV type 16, 18, 31, 33, 35, 39, 45, 51, 52, 56, 58, 59, 66, 68, 69) and 7 low-risk types (HPV type 6, 11, 34, 40, 42, 43, 44) were selected, and genotype-specific probe β -globin for each HPV type possessing an amine group at the 5' terminal of the sequence was prepared for the detection of HPV genotypes. The nucleotide sequence of each probe is set forth in the Sequence List.

Example 2: Preparation of DNA Chip

A DNA chip was prepared as follows: each probe prepared in Example 1 was dissolved in 3X SSC (45mM sodium citrate, 0.45M NaCl, pH 7.0) at a concentration of 200 pmol/ μ l, and spotted onto an aldehyde-derivatized silylated slide (CSS-100, CEL, Houston, TX, USA) to form an array of spots with a size of 150 μ m at 300 μ m of spacing between spots using a microarrayer (GMS 417 Arrayer, TaKaRa, Japan), followed by performing a Schiff's base

reaction under the condition of a 37°C temperature and over 70% humidity, for 4 hours. The slide was washed with 0.2% (w/v) sodium dodecyl sulfate (SDS), and with triple-distilled water. Then the slide was treated with a NaBH₄ solution (0.1g NaBH₄, 30ml phosphate buffered saline (PBS), 10ml ethanol) for 5 minutes to reduce excessive aldehydes not reacted with amine, followed by washing with triple-distilled water and air-drying. Also, β-globin, which functions as a control group for position markers and the hybridization reaction, is affixed to the surface of the solid support.

Example 3: Optimal PCR Amplification

To establish multiplex PCR and amplification of HPV genes from clinical samples, an Optimal HPV PCR was carried out under the following conditions:

3-1: Two-step PCR Amplification

Method 1

To obtain Cy5-labeled amplified DNA samples, PCR was performed with 5 cycles of denaturation for 5 min at 94°C, denaturation for 1 min at 94°C, primer annealing for 2 min at 50°C, and extension for 30 sec at 72°C, followed by 35 cycles of denaturation for 1 min at 94°C, primer annealing for 2 min at 50°C, and extension for 15 sec at 72°C, and then further extension for 2 min at 72°C.

Method 2

Cy5-linked amplified HPV DNA samples were prepared analogously as in Method 1, except for primer annealing for 30 sec at 50°C.

Method 3: Conventional One-step PCR of Prior Art

Cy5-labeled amplified DNA was obtained and compared after 40 cycles of denaturation for 5 min at 94°C, denaturation for 1 min at 94°C, primer annealing for 2 min at 50°C, and extension for 30 sec at 72°C, and then further extension for 2 min at 72°C.

When lanes 4~6, lanes 8~10, and lanes 11~13 in Figure 4 were compared with one another, the conditions in methodology 2 (lanes 4, 8 and 11) were proven to be the most specific and efficient.

3-2: Multiplex PCR Amplification

The compositions and conditions for establishing multiplex PCR amplification of HPV are as follows.

In order to obtain Cy5-labeled amplified DNA samples, multiplex PCR using two-step PCR described in Example 3-1 was performed in a thermocycler (Perkin-Elmer Cetus, CA, USA) with 50 μ l of reaction mixture containing a 2xPCR buffer (50 mM KCl, 4 mM MgCl₂; 10 mM Tris-HCl; pH 8.3); 0.1 μ g of DNA; 2~4 mM MgCl₂; 25 pmol of HPV primers set forth in SEQ ID NO: 43-44; 5 pmol of β -globin primers set forth in SEQ ID NO: 44-45; 40 μ M each of dATP, dCTP, dGTP (Pharmacia); 30 μ M of dTTP (Pharmacia); 0.5 μ M Cy5-dUTP (NEN, USA); 2 units of *Taq* polymerase (TaKaRa, Japan); and optionally BSA and DMSO, with 5 cycles of denaturation for 5 min at 94°C, denaturation for 1 min at 94°C, primer annealing for 2 min at 45~50°C, and extension for 20~30 sec at 72°C, followed by 35 cycles of denaturation for 1

min at 94°C, primer annealing for 2 min at 45~50°C and extension for 10~15 sec at 72°C, and then further extension for 2 min at 72°C.

When lanes 2~5 and lanes 6~9 in Figure 5 were compared, HPV amplification efficiencies of 1xPCR buffer monoplex PCR and of 2xPCR buffer multiplex PCR were similar, and so the 2xPCR buffer multiplex PCR condition was found to be the most specific and efficient.

Example 4: Preparation of Samples.

In order to detect HPV infection in human cervical swabs, DNA was extracted from the said specimen and then purified. To test the adequacy of sample DNA, purified DNA was PCR amplified with β -globin primers set forth in SEQ ID NO: 45 and 46. The DNA samples which revealed β -globin DNA amplification were selected and used for further analyses of HPV DNA.

As HPV DNA standards, plasmid DNA comprising the HPV sequence obtained from the following distributors were used: HPV types 6, 11, 26, 40, 45, 51 and 57 (Dr. Ethel-Michele de Villiers, Angewandte Tumorstudiologie, Deutsches Krebsforschungszentrum, Im Neuenheimer Feld 242, 69009 Heidelberg, Germany); HPV types 35, 44, and 56 (Dr. Attila Lörincz, Vice President, R&D and Scientific Director, Digene Diagnostics, Inc., 2301-B Broadburch Drive, Silver Spring, MD 20904, USA); HPV types 42, 58, 59, 61, 64 and 67 (Dr. Toshihiko Matsukura, Department of Pathology and Laboratory of Pathology, AIDS Research Center, National Institute of Infectious Disease, Tokyo 162, Japan); and HPV types 33, 34, 39, 52, 54, 66 and 70 (Dr. Gérard Orth, Unité Mixte Institut Pasteur/INSERM (U. 190), Institut Pasteur, 25 rue du

Docteur Roux, 75724 Paris Cedex 15, France).

Additionally, DNA extracted and purified from the following cell lines were used as positive controls: SiHa cell line (HPV 16, KCLB 30035, Human squamous carcinoma, cervix), purchased from Korean Cell Line Bank (Seoul National University, College of Medicine, Seoul, Korea).

The selected samples of DNA described above were PCR amplified using primers set forth in SEQ ID NO: 43 and 44.

Example 4-1: Preparation of Positive Control Samples

To obtain Cy5-labeled amplified DNA samples, HPV 16 and HPV 18 DNA purified above were amplified by PCR with primers set forth in SEQ ID NO: 43 and 44. PCR was performed in a thermocycler (Perkin-Elmer Cetus, CA, USA) with 50 μ l of a reaction mixture containing a 1xPCR buffer (50 mM KCl, 4 mM MgCl₂, 10 mM Tris-HCl, pH 8.3); 0.1 μ g of DNA; 4 mM MgCl₂; 25 pmol of each primer, 40 μ M each of dATP, dCTP, and dGTP (Pharmacia); 30 μ M of dTTP (Pharmacia); 0.05 μ M Cy5-dUTP (NEN, USA); and 2 unit of *Taq* polymerase (TaKaRa, Japan) with 5 cycles of denaturation for 5 min at 94°C, denaturation for 1 min at 94°C, primer annealing for 2 min at 50°C, and extension for 30 sec at 72°C, followed by 35 cycles of denaturation for 1 min at 94°C, primer annealing for 2 min at 50°C and extension for 15 sec at 72°C, and then extension for 2 min at 72°C

Example 4-2: Preparation of HPV Standards

Cy5-linked amplified HPV DNA samples were prepared analogously as in Example 4-1, except for employing templates of various HPV plasmids

described above.

Example 4-3: Preparation of Sample DNA from Clinical Samples

Cy5-linked amplified DNA samples were obtained analogously as in Example 4-1, except that DNA obtained from uterine cervical swabs were used
5 as templates, and primers set forth in SEQ ID NO: 43 and 44 were employed as primers.

Example 5: Detection of HPV Infection using DNA Chip

Amplified DNA samples obtained in Example 4 were applied to the DNA chip prepared in Example 2, and hybridization was carried out in a
10 hybridization reaction chamber made up of a Cover slip (GRACE Bio-Labs, USA, PC4L-1.0) with a 100 μl capacity.

As for the quantity of hybridization reaction samples, 10 μl each of amplified product was used for positive controls and plasmid DNA, and a mixture of 10 μl of HPV amplified product and 5 μl of beta-globin amplified
15 product was used for DNA obtained from cervical swabs. The said reaction samples were denatured by adding 3N NaOH solution (10% v/v) and standing them for 5 min at room temperature, and neutralized by adding 1 M Tris-HCl (pH 7.2, 5% v/v) followed by 3N HCl (10% v/v) and cooling them for 5 min on ice. The samples were then mixed with a hybridization solution made up of a
20 6X SSPE (saline-sodium phosphate-EDTA buffer, Sigma Chemical Co., St. Louis, MO, USA) and 0.2% SDS (sodium dodecyl sulfate), and applied onto the DNA chip. The hybridization reaction was carried out for 2 hours at 40°C, followed by washing with a 3X SSPE for 2 min and a 1X SSPE for 2 min, and

air-drying at room temperature or spin-dryer drying.

The DNA chip hybridized with sample DNA was analyzed for fluorescent signals (extinction 650 nm, emission 668 nm) by using a confocal laser scanner (GSI Lumonics, Germany) (see: Figure 1, Figures 2a-2l and
5 Figures 3a-3j).

Figure 1 is a schematic representation of the type and position of the probes on the DNA chip: each number indicates each HPV probe, circles(○) indicate HPV and beta-globin probe-affixed positions, and closed circles(●) indicate background markers to verify proper performance of the hybridization
10 reaction and position markers for locating probes. Figures 2a-2e are photographs showing the results of high-risk group HPV DNA analyses using HPV plasmids and cervical cancer cell lines (HPV 16, 18, 39, 58 and 69), and Figures 2g-2i are photographs showing the results of low-risk group HPV DNA analyses using HPV plasmids (HPV 6, 11 and 40). As shown in Figures 2a-2e
15 and Figures 2g-2i, hybridization signals produced by the amplified DNA of HPV plasmid standards and HPV positive controls (cervical cancer cell lines) were observed clearly on the corresponding probes without significant cross-hybridization.

Example 6: Detection of HPV Infection in Clinical Samples using DNA

20 Chip

In order to examine the accuracy and efficiency of diagnosis by the DNA chip of the invention, clinical samples were PCR amplified with primers comprising nucleotide sequences set forth in SEQ ID NO: 43-46, and then, for proper samples, a diagnostic procedure using the DNA chip was performed to

detect HPV infection as well as to determine the type of infection.

DNA isolated from 213 uterine cervix specimens were amplified using the method described in Example 4-3, and analyzed for HPV infection by using the DNA chip of the invention. Detection of HPV included in the high-risk group was carried out using a Hybrid Capture II method provided by the Digene Diagnostics company. The results of the two methods were compared (see: Figures 2a-2l, Figures 3a-3j, and Table 3). Figures 2a-2l and Figures 3a-3j are photographs showing Examples of the results of DNA chip analyses of cervical swab specimens for HPV infection. As shown in the Figures, detailed diagnoses of HPV infection in the clinical samples via accurate detection and genotyping of the infecting HPV were successfully accomplished by using the DNA chip of the invention. The results of the DNA chip analyses and Hybrid Capture II analysis are shown in Table 3:

[Table 3]

Hybrid Capture II	DNA Chip of the Present Invention			Percent (%)
	Group	HPV Type	The Number of Single-infected Samples (The Number of Whole Infected samples)	
Positive 149	High-risk Group	HPV 16	31 *(66)	20.8 (44.3)
		HPV 18	1 (18)	0.7 (12.1)
		HPV 31	2 (3)	1.4 (2)
		HPV 33	2 (6)	1.4 (4)
		HPV 35	2 (6)	1.4 (4)
		HPV 39	1 (3)	0.7 (2)
		HPV 45	- (1)	- (0.7)
		HPV 51	6 (9)	4 (6)
		HPV 52	10 (14)	6.7 (9.4)
		HPV 56	15 (27)	10.1 (18.1)
		HPV 58	7 (13)	4.7 (8.7)
		HPV 59	2 (7)	1.4 (4.7)
		HPV 66	1	0.7
		HPV 68	2 (6)	1.4 (4)
		HPV 69	-	-
	Low-risk Group	HPV 6	- (1)	- (0.7)
		HPV 11	1 (2)	0.7 (1.4)
		HPV 34	1 (2)	0.7 (1.4)
		HPV 40	1 (2)	0.7 (1.4)
		HPV 42	-	
		HPV 43	1	0.7
		HPV 44	- (1)	- (0.7)
	Other type		4	2.7
Negative 51	Negative 50	HPV 18 1		

*1(): the number of samples comprising multiple-infected samples

In Table 1, "-" indicates absence of sample infected with HPV in the corresponding item, and percent was calculated by {(the number of single-infected samples or total number of infected samples) / 149} x 100.

5

[Table 4]

Specificity and Sensitivity of DNA chip according to the invention

DNA Chip Of The Invention	Hybrid Capture II		
	Positive	Negative	Sum
Positive	162	1*	163
Negative	0	50	50
Sum	162	51	213
Relative Sensitivity			100%
Relative Specificity			98% (95%CI=94-100)
Similarity			98.6% (95%CI=99-100)
Copy Value(κ)			0.99 (95%CI=0.96-1.00)

As shown in Table 4, DNA chip analysis according to the invention was reported to have 100% relative sensitivity and 98% relative specificity as compared with the Hybrid Capture assay. The above information indicates that diagnosis of HPV infection using the genotyping kit of the invention is superior in many aspects to the conventional methods employed for the same purpose.

In Tables 3 and 4, all samples which were confirmed as positive by Hybrid Capture assay were tested or diagnosed with the DNA chip of the invention, and one of samples which were confirmed as positive by Hybrid Capture assay was detected to be HPV 18 type. As a result of sequencing to determine whether it was a false-positive, it was confirmed that it was HPV 18.

While the Hybrid Capture assay can only diagnosis HPV infection, the DNA chip analysis can detect HPV infection and identify the types of infecting HPV in a fast and accurate manner. The DNA chip analysis according to the invention is so highly sensitive that it can detect the samples confirmed to be false-negative through the Hybrid Capture assay. As well-known to those skilled in the art, HIV16 type shows the highest frequency in HIV infection, followed byHIV 56, 58 and 52 types in contrast to the Western tendency with HPV 18 being the second frequent type.

As clearly illustrated and demonstrated above, the present invention provides a genotyping kit comprising said probes (SEQ ID NO: 1 to 30) for identifying genotypes of HPV from clinical samples of infected patients and a method for diagnosis of HPV infection by genotyping the infecting virus using the said genotyping kit.

Comparative Example 1

As shown in Table 6, efficiency of the probes according to the invention was compared with that of the conventional probes for HPV 16, 34, 56, and 59, and a DNA chip to confirm efficiencies of additional probes for HPV 43, 68, and 69 was described. A test was carried out analogously as in Example 6, except for employing the DNA chip described above.

As a result of analysis of control DNA for each type in Figures 7-8, conventional probes were found to pose non-specificity (false positive) and low-sensitivity (false negative).

Comparative Example 2: DNA Chip Test

A test was carried out analogously as in Example 6, except for

employing the conventional probes including probes set forth in SEQ ID NO: 31-41 and 47-54 and using the DNA chip of the invention. Sequencing was performed using the conventional sequencing method.

Results of analysis using the conventional DNA chip art and the present DNA chip, results of sequencing via PCR for identification, and results of analysis using the conventional probes and the present probes are shown in Tables 5a and 5b. It was found that in the frequency of false negative and false positive, the genotyping kit using the conventional DNA chip is higher than the genotyping kit using the DNA chip of the invention.

The present invention provides a genotyping kit for diagnosing patients infected with human papillomavirus (HPV), which includes the said probes, and a method for genotyping of HPV DNA isolated from patients using the said genotyping kit in an accurate and fast manner.

[Table 5a]

Sample	Conventional HPV DNA Chip	DNA Chip Of The Invention	Sequencing	Conventional HPV DNA Chip	
				False Positive	False Negative
1	OT	16,68	16,68		16, 68
2	35	51	51	35	51
3	16	OT	OT	16	
4	59	59,68	59,68		68
5	40, 58	51	51	40, 58	51
6	16, 18, 33, 35	16,18,33	16,18,33	35	
7	33	16,33	16,33		16
8	16	16,68	16,68		68
9	58	16,18	16,18	58	16, 18
10	16	56	56	16	56
11	16	56,58	56,58	16	56, 58
12	16	56	56	16	56

13	35	68	68	35	
14	16	16, 69	16, 69		69
15	35, 52	52	52	35	52
16	16	18, 56	18, 56	16	18, 52
17	35	52	52	35	52
18	35	51	51	35	51
19	58	40	40	58	40
20	59	52	52	59	52
21	59	16	16	59	16
22	35	16, 18	16, 18	35	16, 18
23	35	16, 18	16, 18	35	16, 18
24	35	43	43	35	43
25	16	52	52	16	52
26	35, 16	35	35	16	
27	58	31, 39, 58	31, 39, 58		31, 39
28	31, 59	31	31	59	31
29	31, 16	31	31	16, 35	31
30	35, 16	31	31	16, 35	31
31	35, 58	35, 58	35, 58		
32	35, 33	33	33	35	33

[Table 5b]

33	16, 35	16	16	35	
34	52, 40	52, 66	52, 66	40	66
35	58	66, 58	66, 58		66
36	16, 56	16	16	56	
37	58, 34	58	58	34	
38	16	16, 43	16, 43		43
39	31, 35	31	31	35	
40	33, 16	33	33	16	
41	16	18	18	16	
42	16, 40	neg	neg	16, 40	
43	16, 56	neg	neg	16, 56	
44	16, 59	16	16	59	
45	35	35, 66	35, 66		66
46	16, 40	16	16	40	
47	34, 58	58	58	34	
48	6, 18	6, 18, 69	6, 18, 69		69
49	58	51	51	58	51
50	59	11	11	59	11

The HPV genotyping kit of the invention is an implement that can detect HPV infection in a simple and accurate manner, as well as identify the types of infecting HPV, and therefore it may contribute to early diagnosis, prevention, and treatment of cervical cancer.

While the present invention has been described in detail with reference to the preferred embodiments, those skilled in the art will appreciate that various modifications and substitutions can be made thereto without departing from the spirit and scope of the present invention as set forth in the appended claims.

WHAT IS CLAIMED IS:

1. A probe comprising a nucleotide sequence which can complementarily bind to DNA Papillomavirus (HPV and is selected from the group consisting of oligonucleotides having nucleotide sequences set forth in
5 SEQ ID NO: 1 to 30 and oligonucleotides having nucleotide sequences complementary to the said oligonucleotide.
2. A DNA chip for genotyping HPV comprising at least one HPV probes selected from the group consisting of probes according to Claim 1.
3. The DNA chip according to claim 2, wherein the DNA chip further
10 comprises one or more HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.
4. The DNA chip according to claim 3, wherein the DNA chip comprises:
15
 - (i) one or more HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 1 to 20 and nucleotide sequences complementary to the said nucleotide sequences;
 - (ii) 8 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide
20 sequences complementary to the said nucleotide sequences; and
 - (iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.
5. The DNA chip according to Claim 4, wherein the DNA chip

comprises:

(i) 9 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 4 to 10;

5 (ii) 8 HPV probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide sequences complementary to the said nucleotide sequences; and

(iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

10 6. The DNA chip according to Claim 5, wherein the DNA chip comprises:

(i) 3 HPV probes having nucleotide sequences set forth in SEQ ID NO: 4, 15, and 16;

15 (ii) 8 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide sequences complementary to the said nucleotide sequences; and

(iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

20 7. An HPV genotyping kit which comprises:

(i) a DNA chip for HPV genotyping according to any one of Claims 2 to 6;

(ii) primers for amplifying sample DNA by PCR; and

(iii) means for labeling amplified DNA hybridized with the said DNA

chip.

8. The HPV genotyping kit according to Claim 7, wherein the DNA chip further comprises position markers to locate probes.

9. The HPV genotyping kit according to Claim 8, wherein the position
5 markers are selected from the group consisting of β -globin, actin, and glyceraldehyde-3-phosphate dehydrogenase gene.

10. The HPV genotyping kit according to Claim 7, wherein the means for labeling are selected from the group consisting of Cy5, Cy3, biotin-binding material, EDANS (5-(2'-aminoethyl)amino-1-naphthalene sulfuric acid),
10 tetramethylrhodamine (TMR), tetramethylrhodamine isocyanate (TMRITC), x-rhodamine, and Texas red.

11. The HPV genotyping kit according to Claim 7, wherein the amplifying of sample DNA by PCR is two-step amplification in which a chain-extension step, that is, the first amplification step and the second amplification
15 step, are carried out for 20-30 seconds and 10-15 seconds, respectively.

12. The HPV genotyping kit according to Claim 9 or 11, wherein the amplifying of sample DNA by PCR is carried out by simultaneously adding primers for amplifying HPV and primers for amplifying β -globin.

13. The HPV genotyping kit according to Claim 7, wherein the
20 primers comprise primers for amplifying β -globin which has a nucleotide sequence set forth in SEQ ID NO: 45 or 46.

14. A primer for amplifying β -globin which has a nucleotide sequence set forth in SEQ ID NO: 45 or 46.

AMENDED CLAIMS

[received by the International Bureau on 20 November 2002 (20.11.02);
original claims 1, 3, 5 and 12 amended: remaining claims unchanged (3 pages)]

1. (Amended) A probe comprising a nucleotide sequence which can complementarily bind to DNA of Human Papillomavirus (HPV) and is selected from the group consisting of oligonucleotides having nucleotide sequences set forth in SEQ ID NO: 1, SEQ ID NO: 3 to 28, and SEQ ID NO: 30, and oligonucleotides having nucleotide sequences complementary to the said oligonucleotide.

2. A DNA chip for genotyping HPV comprising at least one HPV probe selected from the group consisting of probes according to Claim 1.

10 3. (Amended) The DNA chip according to claim 2, wherein the DNA chip further comprises one or more HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 2, SEQ ID NO: 29, and SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

15 4. The DNA chip according to claim 3, wherein the DNA chip comprises:

(i) one or more HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 1 to 20, and nucleotide sequences complementary to the said nucleotide sequences;

20 (ii) 8 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide sequences complementary to the said nucleotide sequences; and

(iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID

NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

5. (Amended) The DNA chip according to Claim 4, wherein the DNA chip comprises:

5 (i) 9 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 1 to 20;

(ii) 8 HPV probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide sequences complementary to the said nucleotide sequences; and

10 (iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

6. The DNA chip according to Claim 5, wherein the DNA chip comprises:

15 (i) 3 HPV probes having nucleotide sequences set forth in SEQ ID NO: 4, 15, and 16;

(ii) 8 HPV probes selected from the group consisting of probes having nucleotide sequences set forth in SEQ ID NO: 21 to 30 and nucleotide sequences complementary to the said nucleotide sequences; and

20 (iii) 11 HPV probes having nucleotide sequences set forth in SEQ ID NO: 31 to 41 and nucleotide sequences complementary to the said nucleotide sequences.

7. An HPV genotyping kit which comprises:

(i) a DNA chip for HPV genotyping according to any one of Claims 2 to 6;

(ii) primers for amplifying sample DNA by PCR; and

(iii) means for labeling amplified DNA hybridized with the said DNA
5 chip.

8. The HPV genotyping kit according to Claim 7, wherein the DNA chip further comprises position markers to locate probes.

9. The HPV genotyping kit according to Claim 8, wherein the position markers are selected from the group consisting of β -globin, actin, and
10 glyceraldehyde-3-phosphate dehydrogenase gene.

10. The HPV genotyping kit according to Claim 7, wherein the means for labeling are selected from the group consisting of Cy5, Cy3, biotin-binding material, EDANS (5-(2'-aminoethyl)amino-1-naphthalene sulfuric acid), tetramethylrhodamine (TMR), tetramethylrhodamine isocyanate (TMRITC), x-
15 rhodamine, and Texas red.

11. The HPV genotyping kit according to Claim 7, wherein the amplifying of sample DNA by PCR is two-step amplification in which a chain-extension step, that is, the first amplification step and the second amplification step, are carried out for 20-30 seconds and 10-15 seconds, respectively.

20 12. (Amended) The HPV genotyping kit according to Claim 9, wherein the amplifying of sample DNA by PCR is carried out by simultaneously adding primers for amplifying HPV and primers for amplifying β -globin.

13. The HPV genotyping kit according to Claim 7, wherein the primers

STATEMENT UNDER ARTICLE 19 (1)

The Applicant, who received the International Search Report relating to the above-identified International Application transmitted on June 25, 2002, hereby files amendment under Article 19 as in the attached sheet.

Therefore, the Applicant hereby amends claims 1, 3, 5, and 12 as originally filed at the international filing date. Please substitute the enclosed new set of claims for all the claims as originally filed.

FIG. 1A

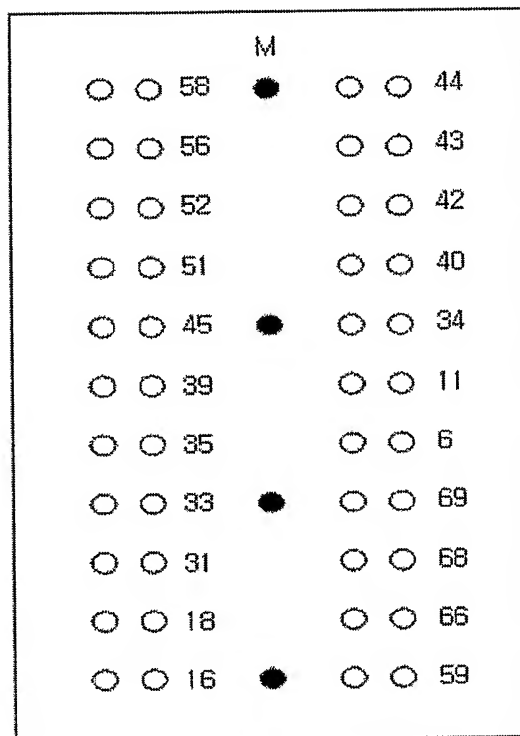


FIG.1B

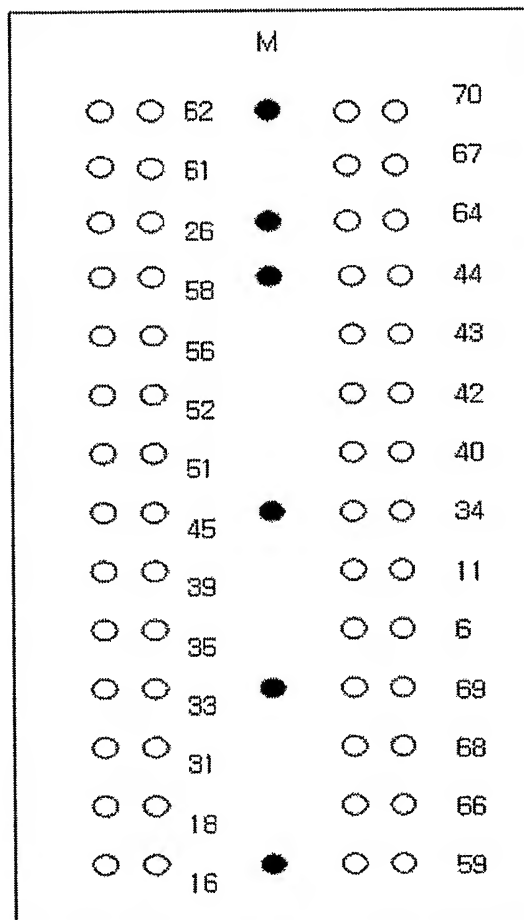


FIG.2A

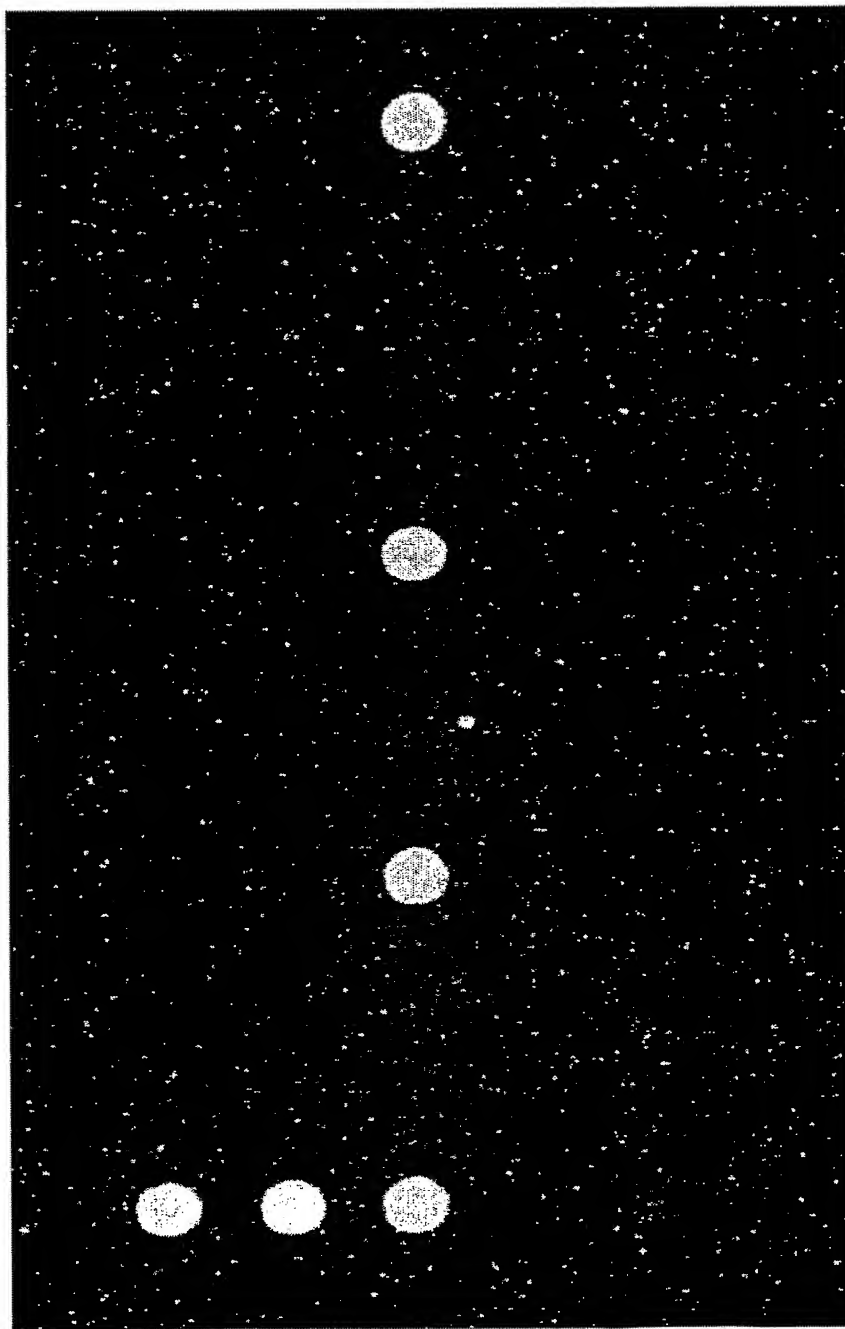


FIG.2B

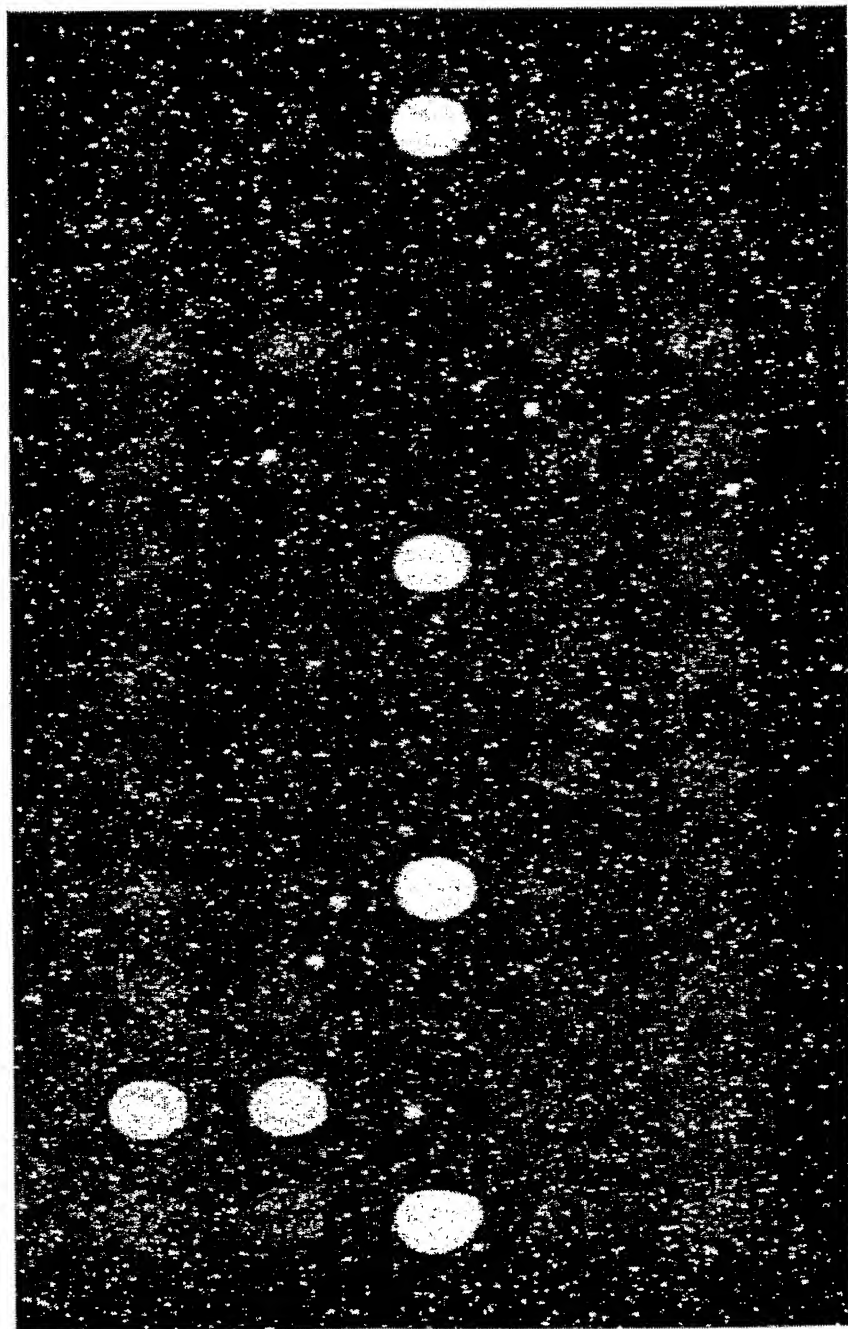


FIG.2C

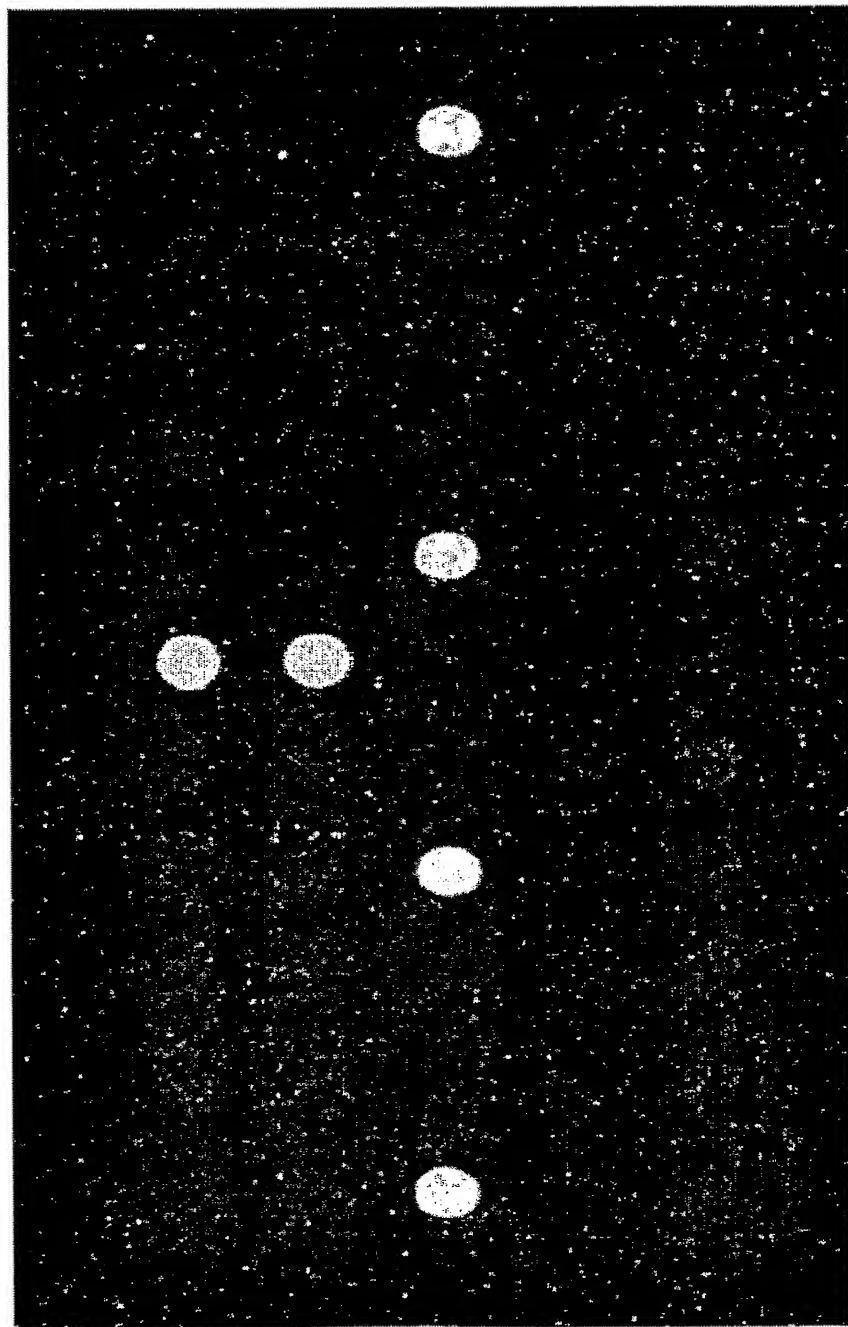


FIG.2D

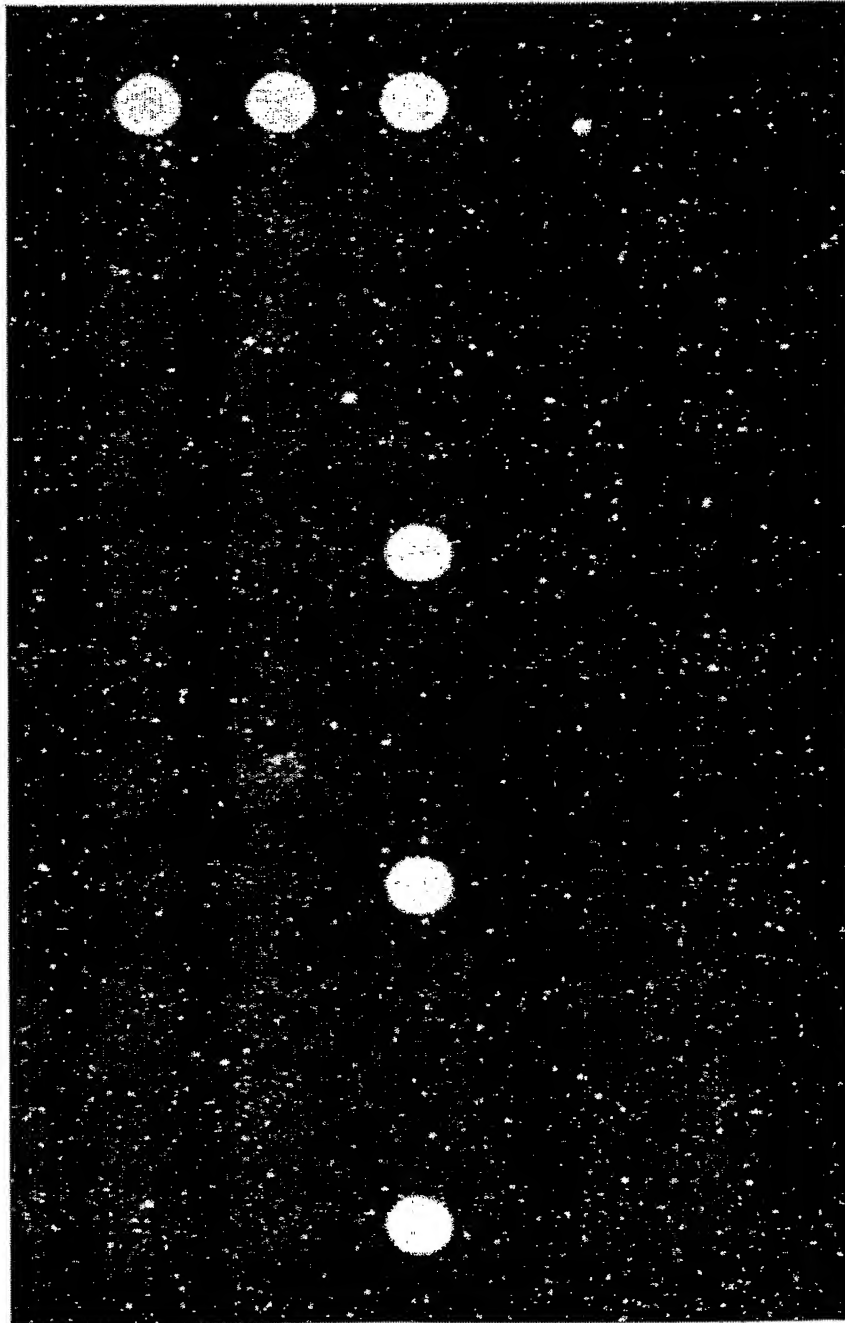


FIG.2E

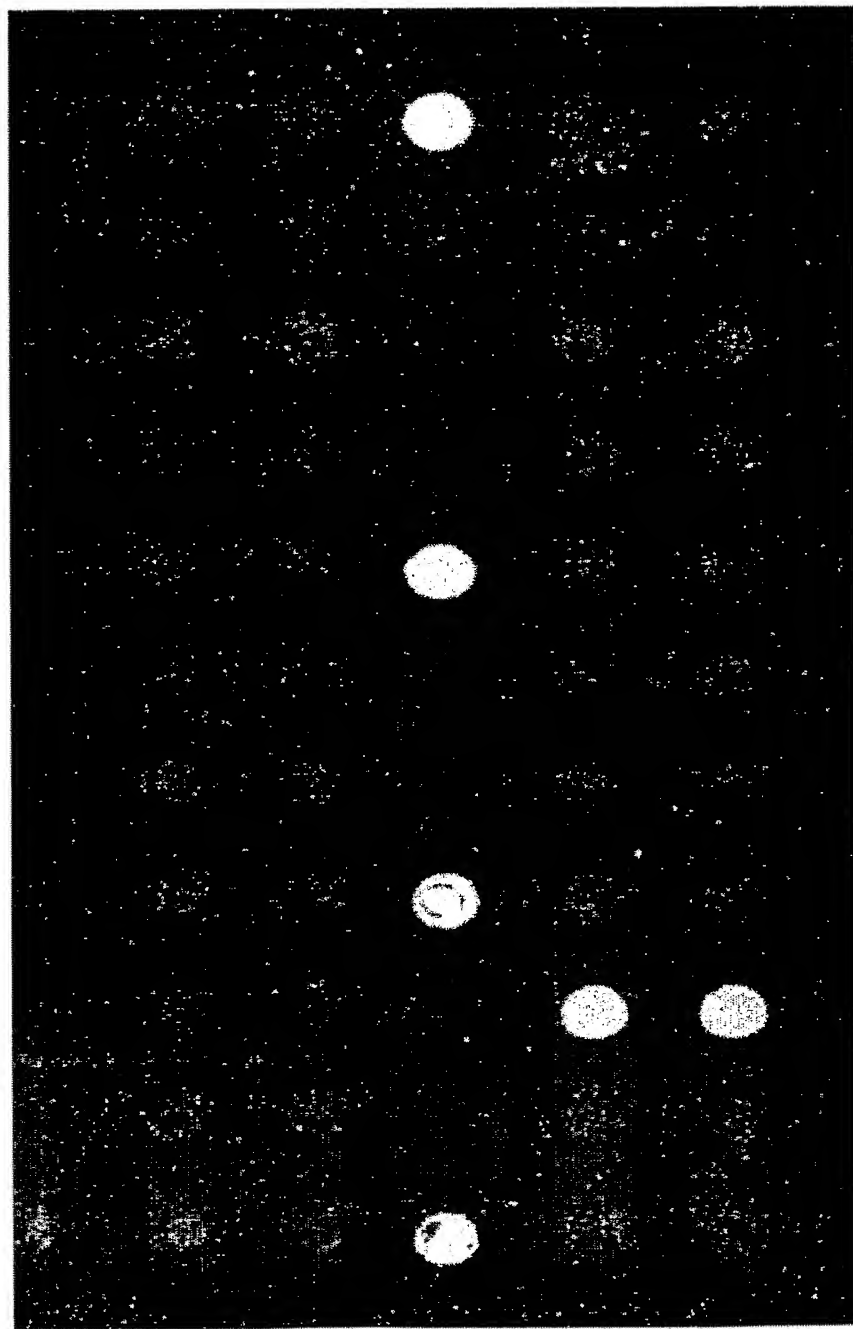


FIG.2F

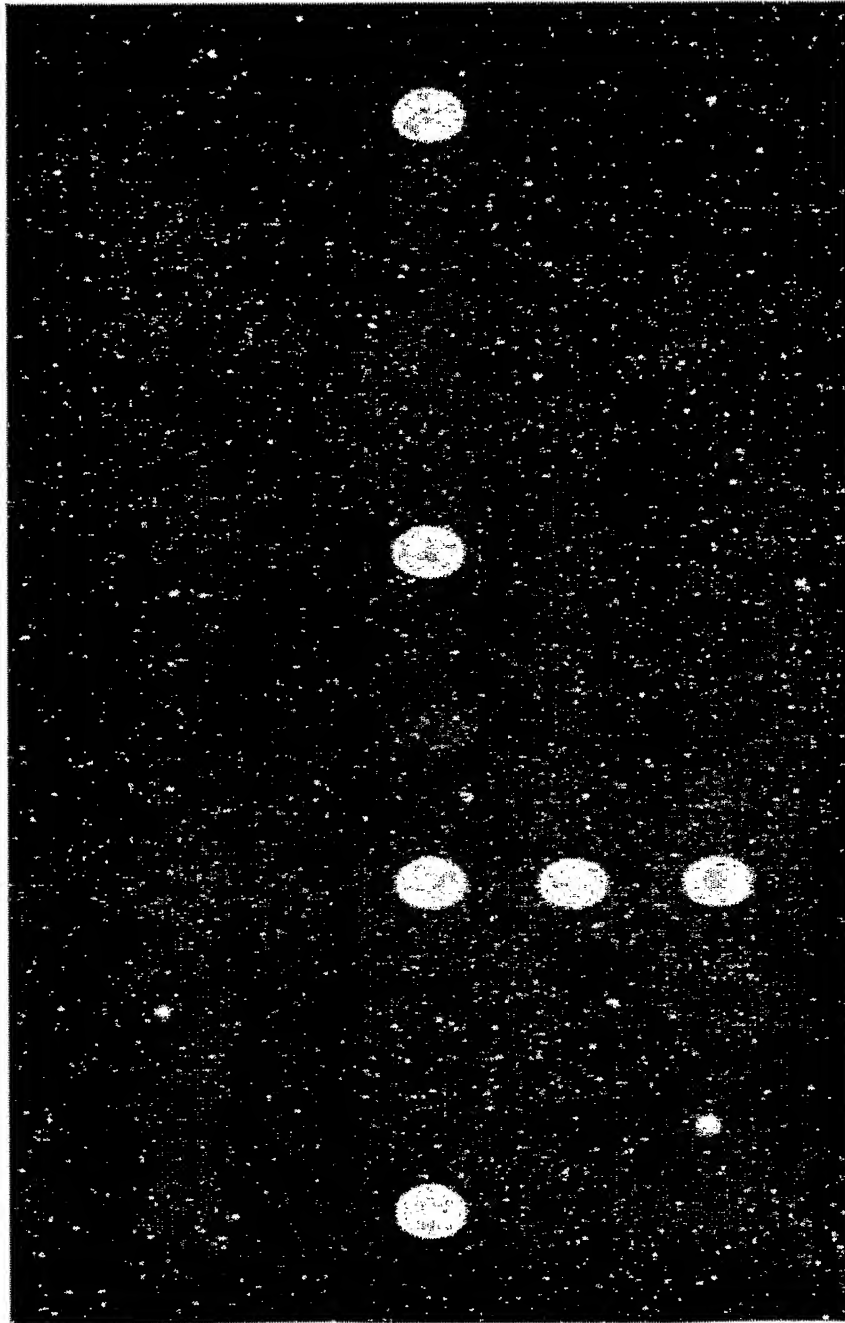


FIG.2G



FIG.2H

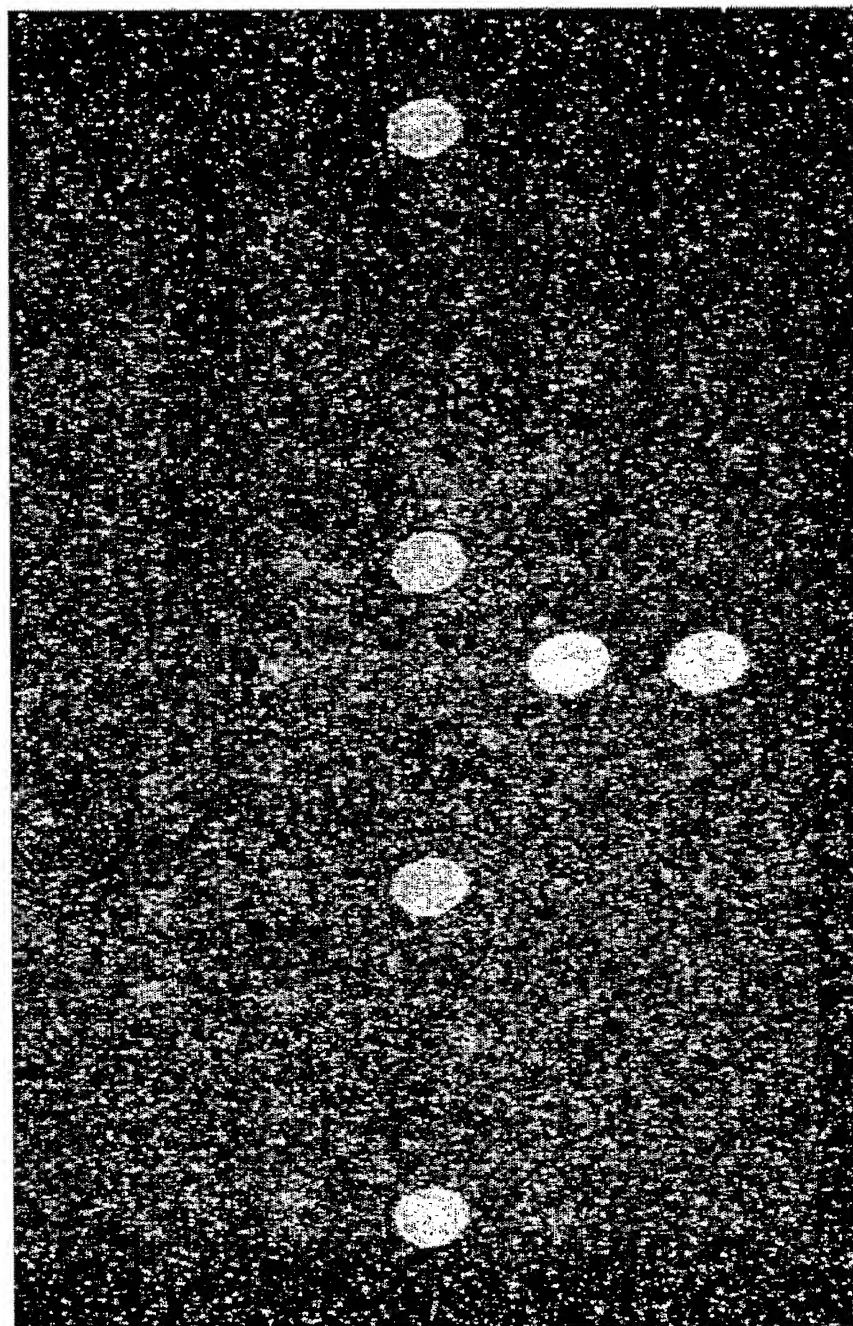


FIG.2I



FIG.2J

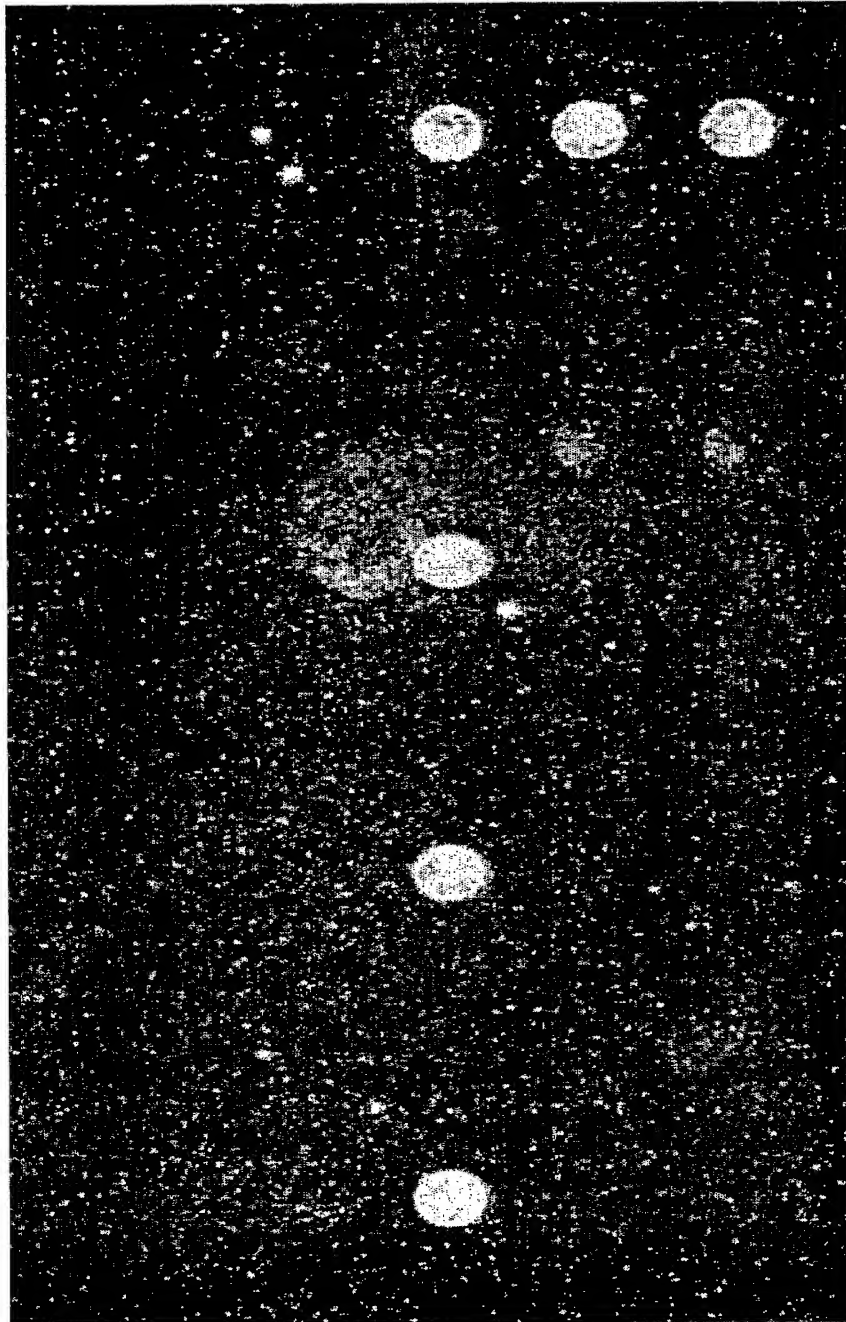


FIG.2K

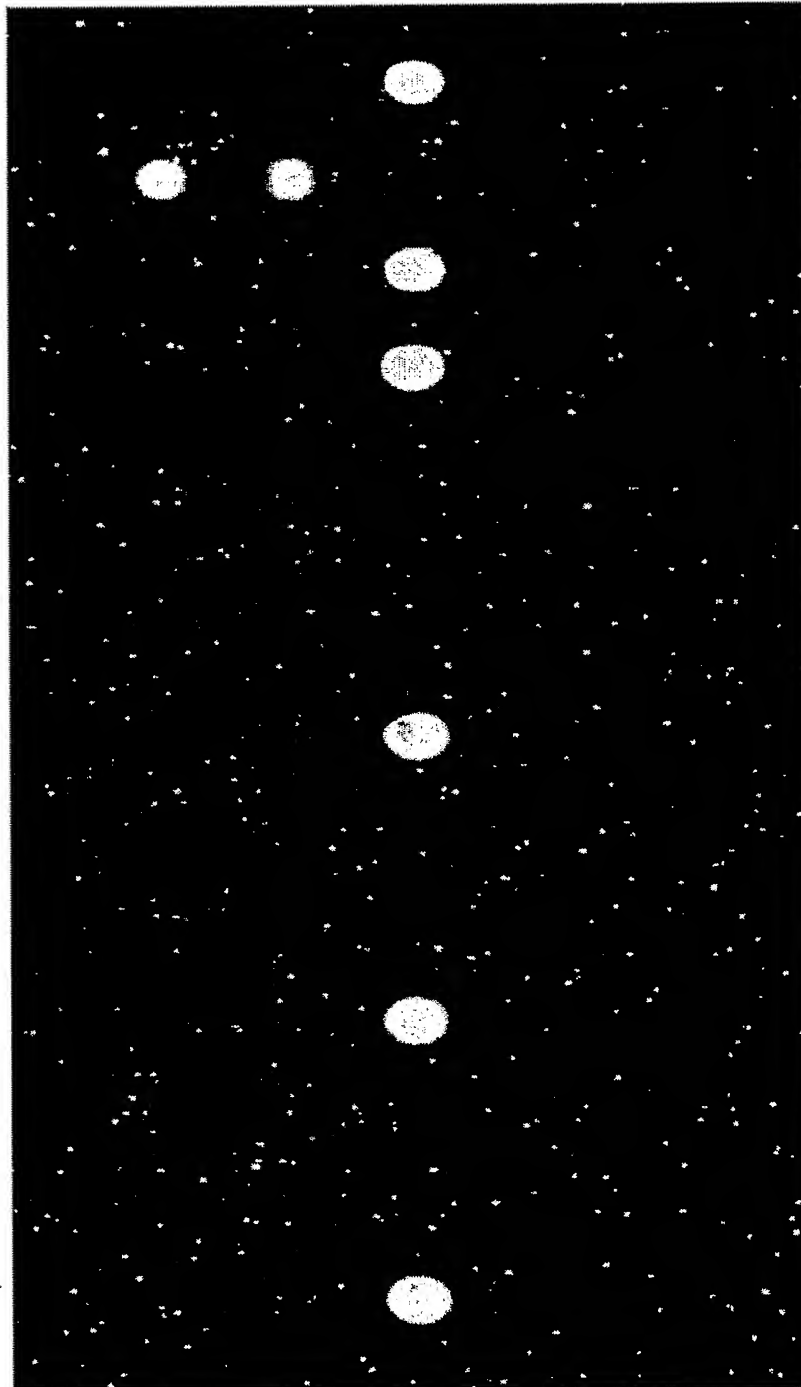


FIG.2L

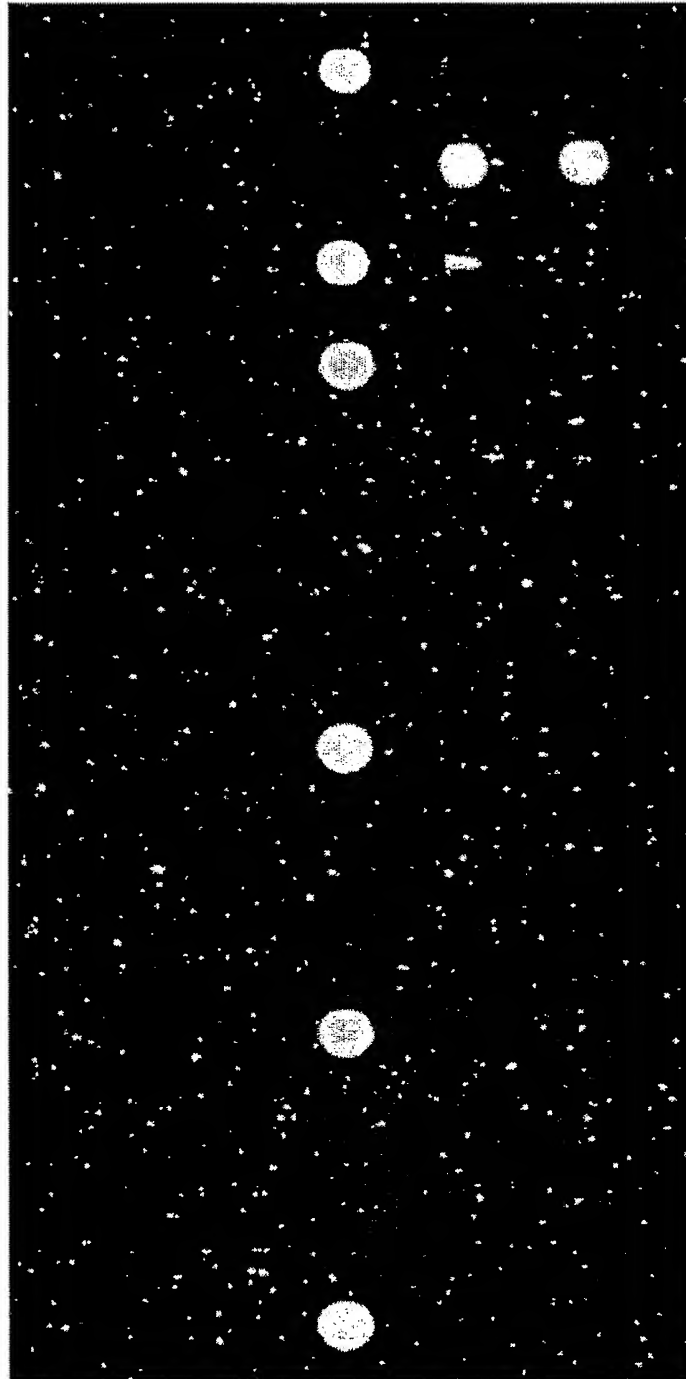


FIG.3A

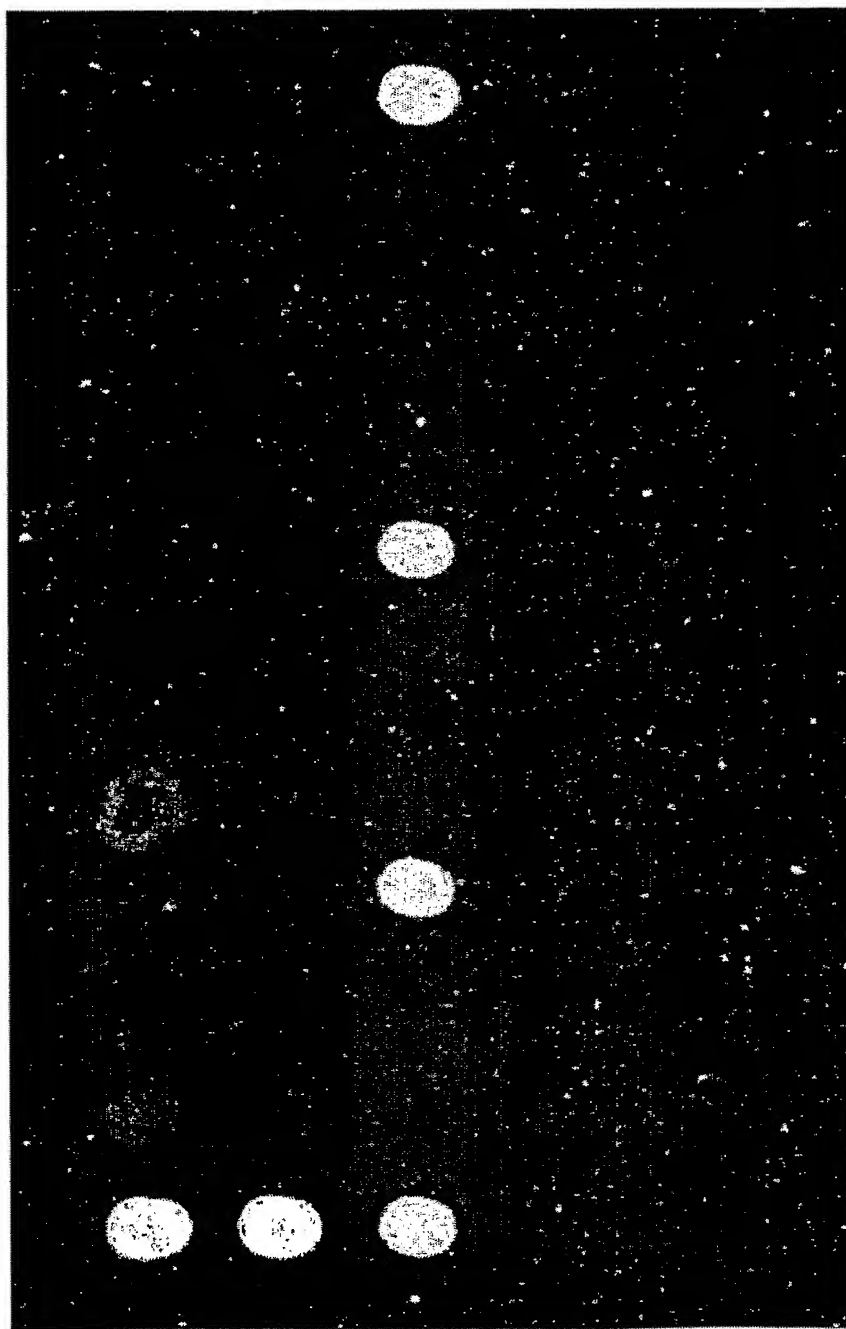


FIG.3B

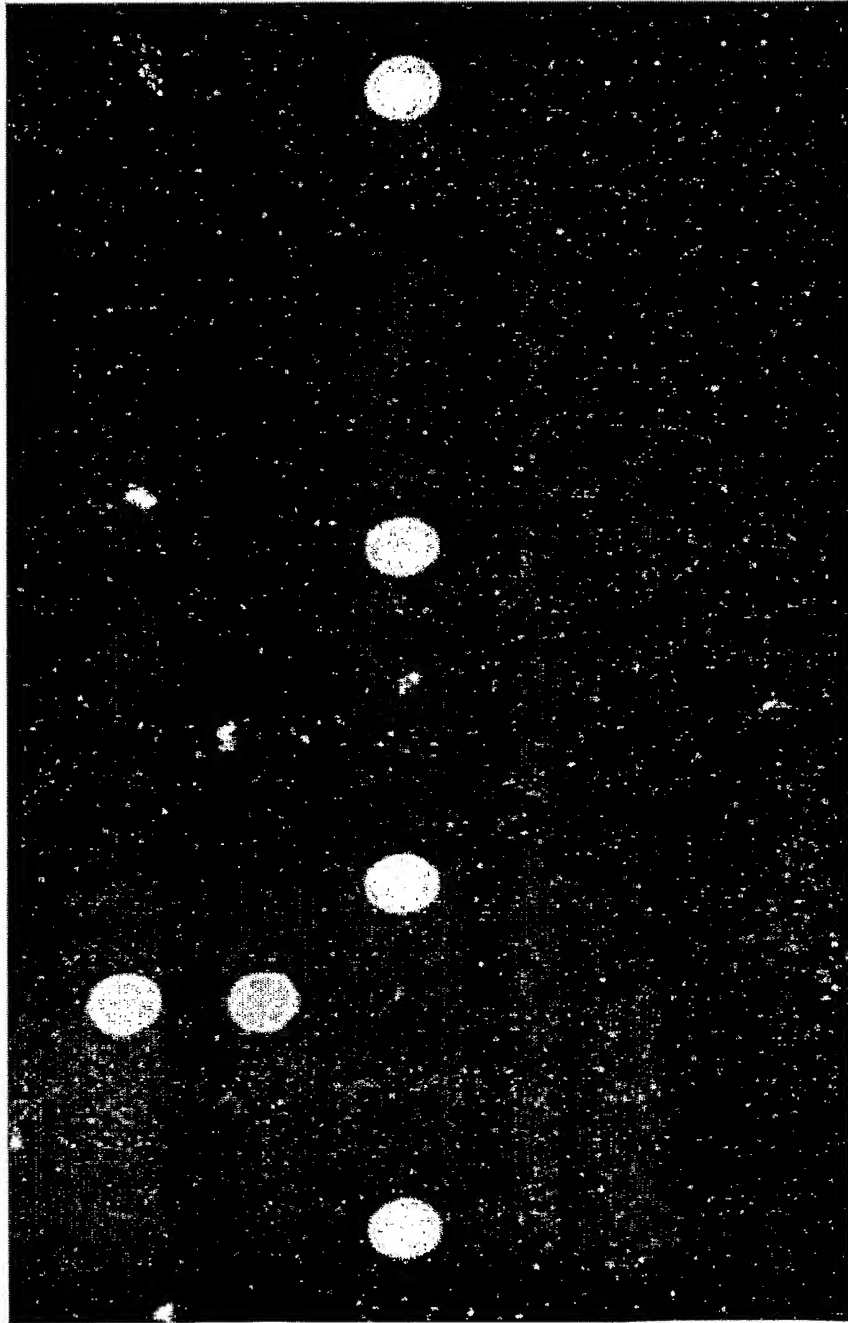


FIG.3C

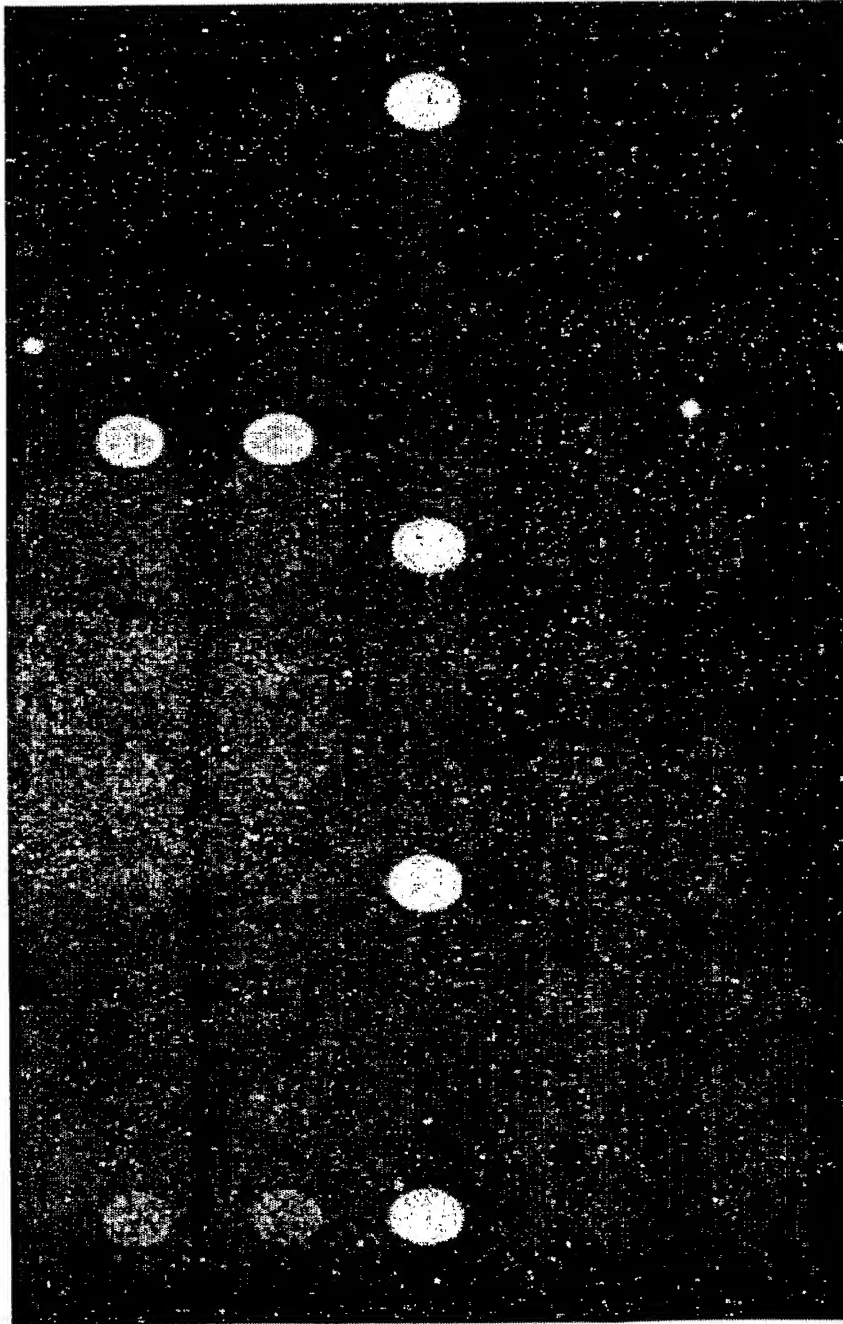


FIG.3D



FIG.3E

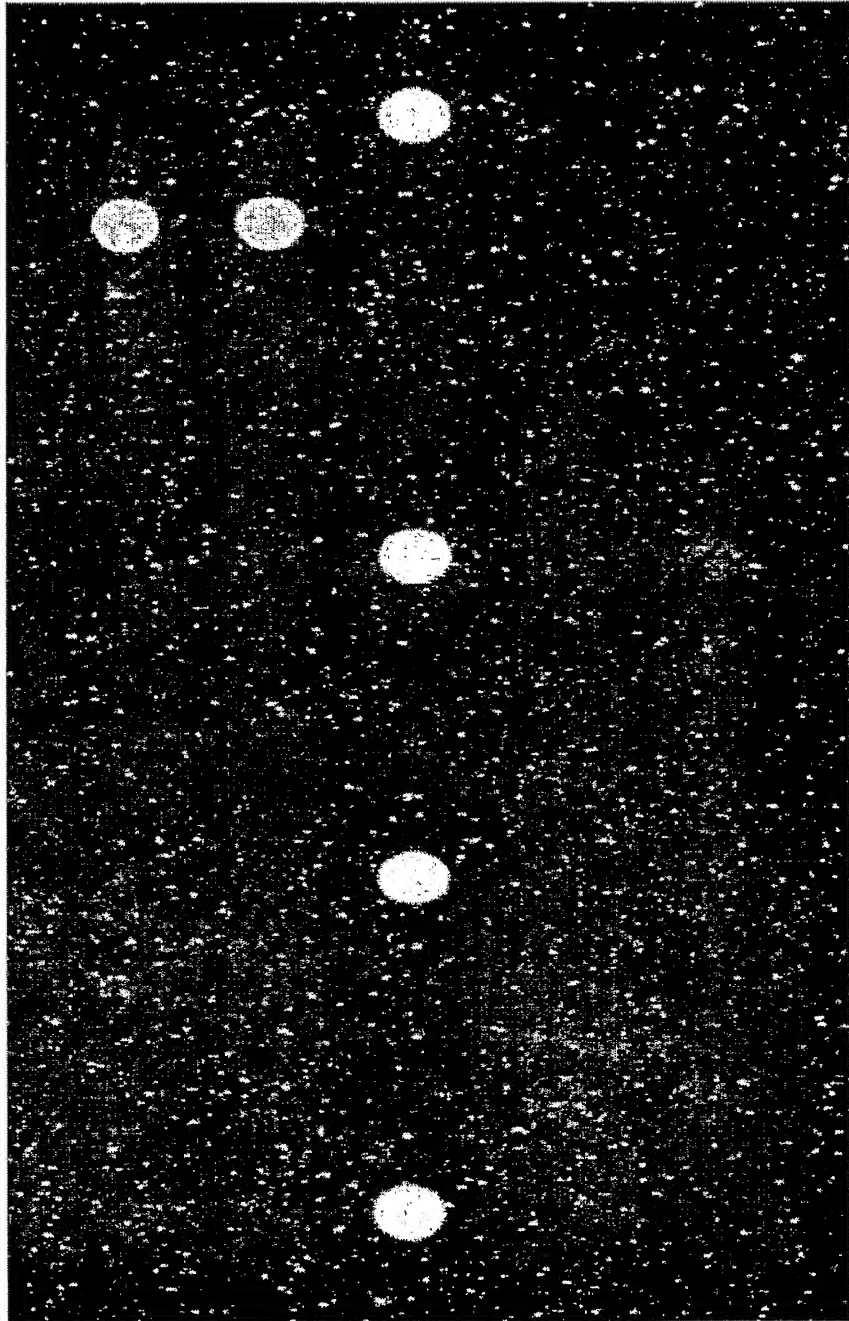


FIG.3F

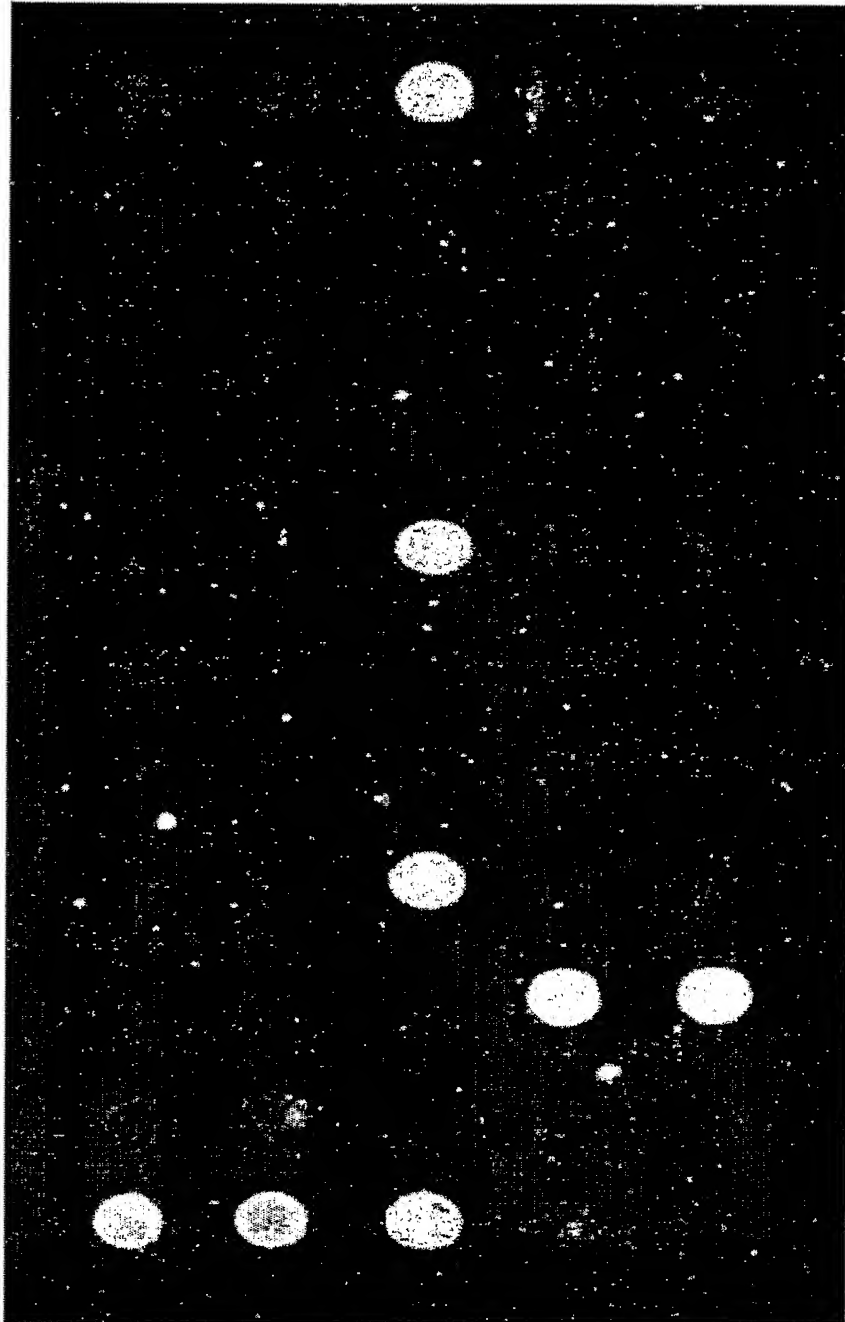


FIG.3G

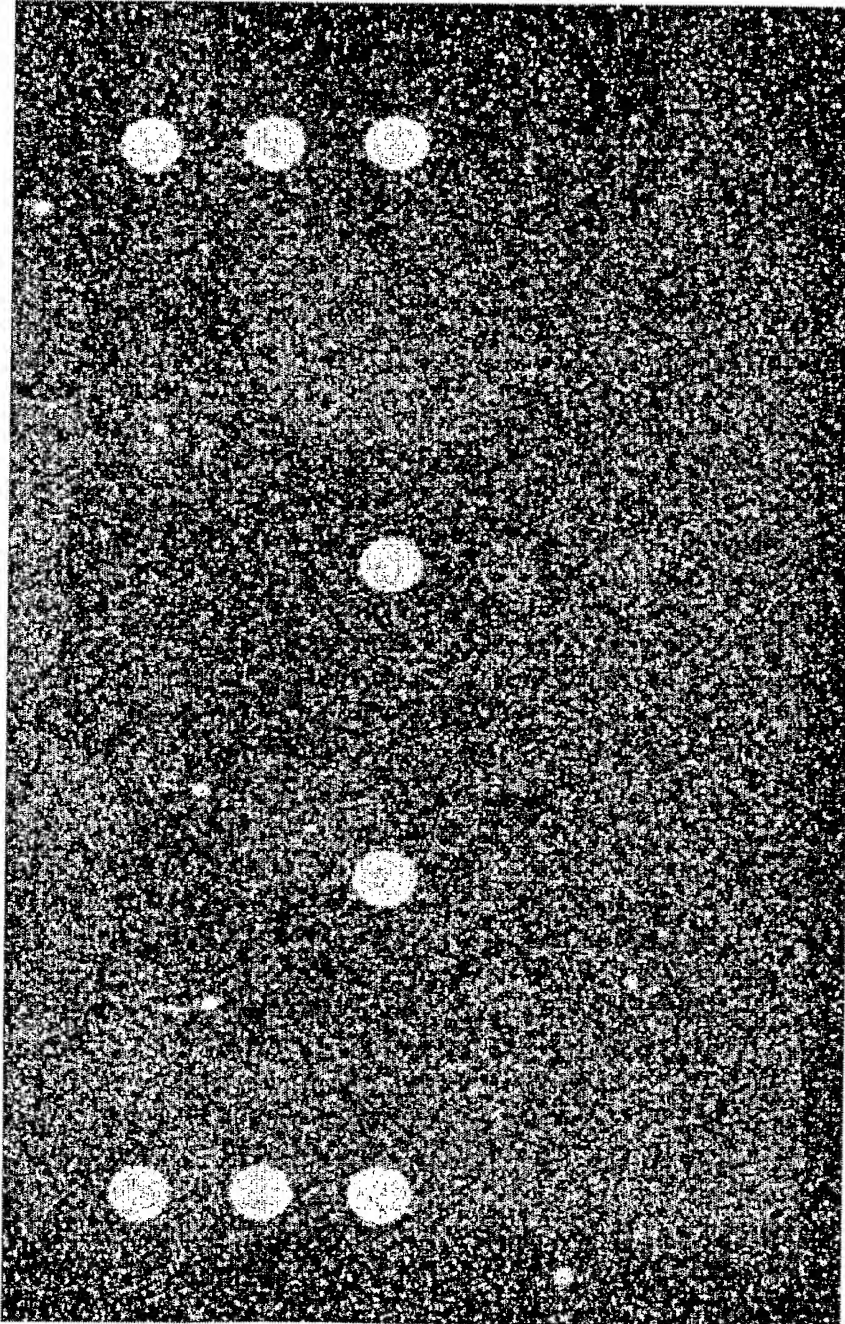


FIG.3H

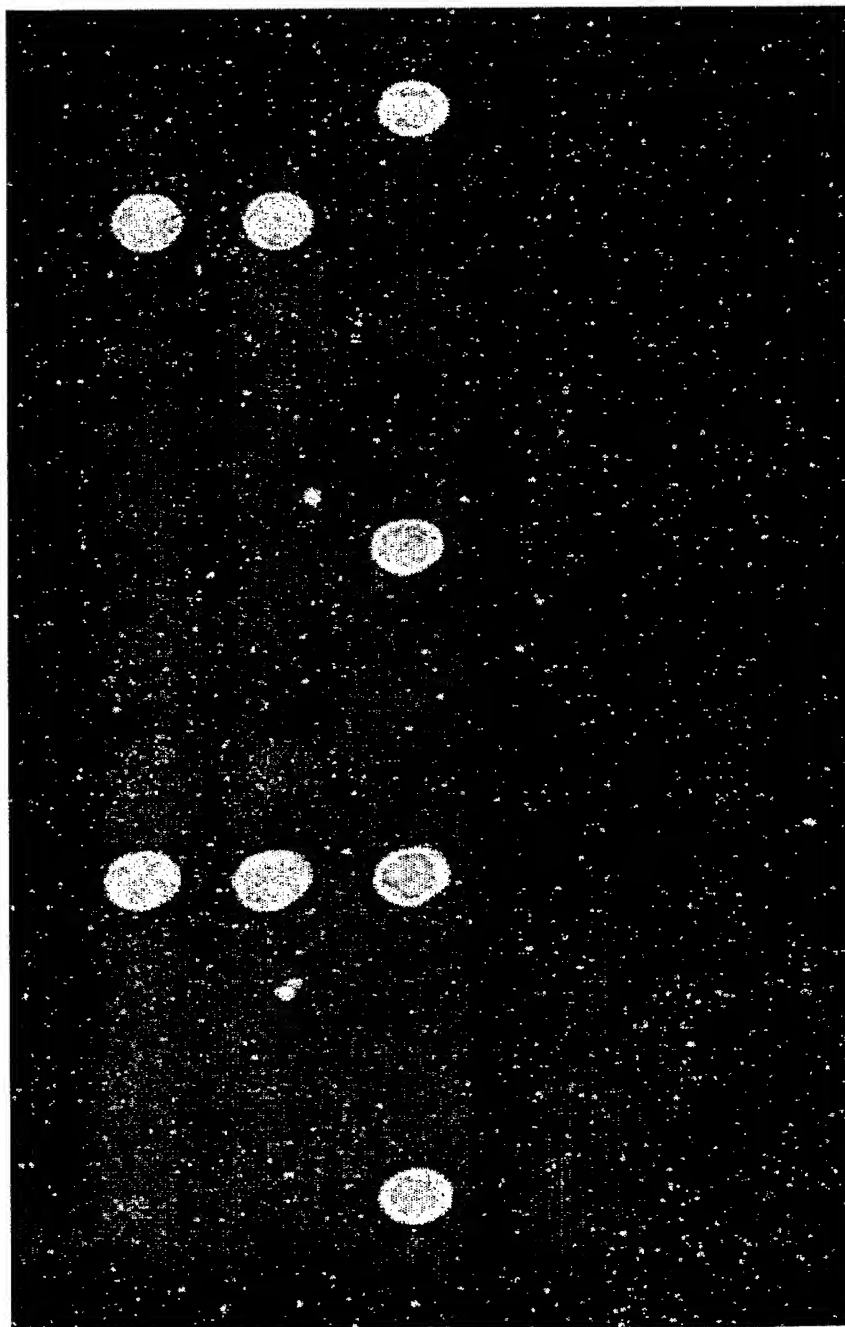


FIG.3I



FIG.3J

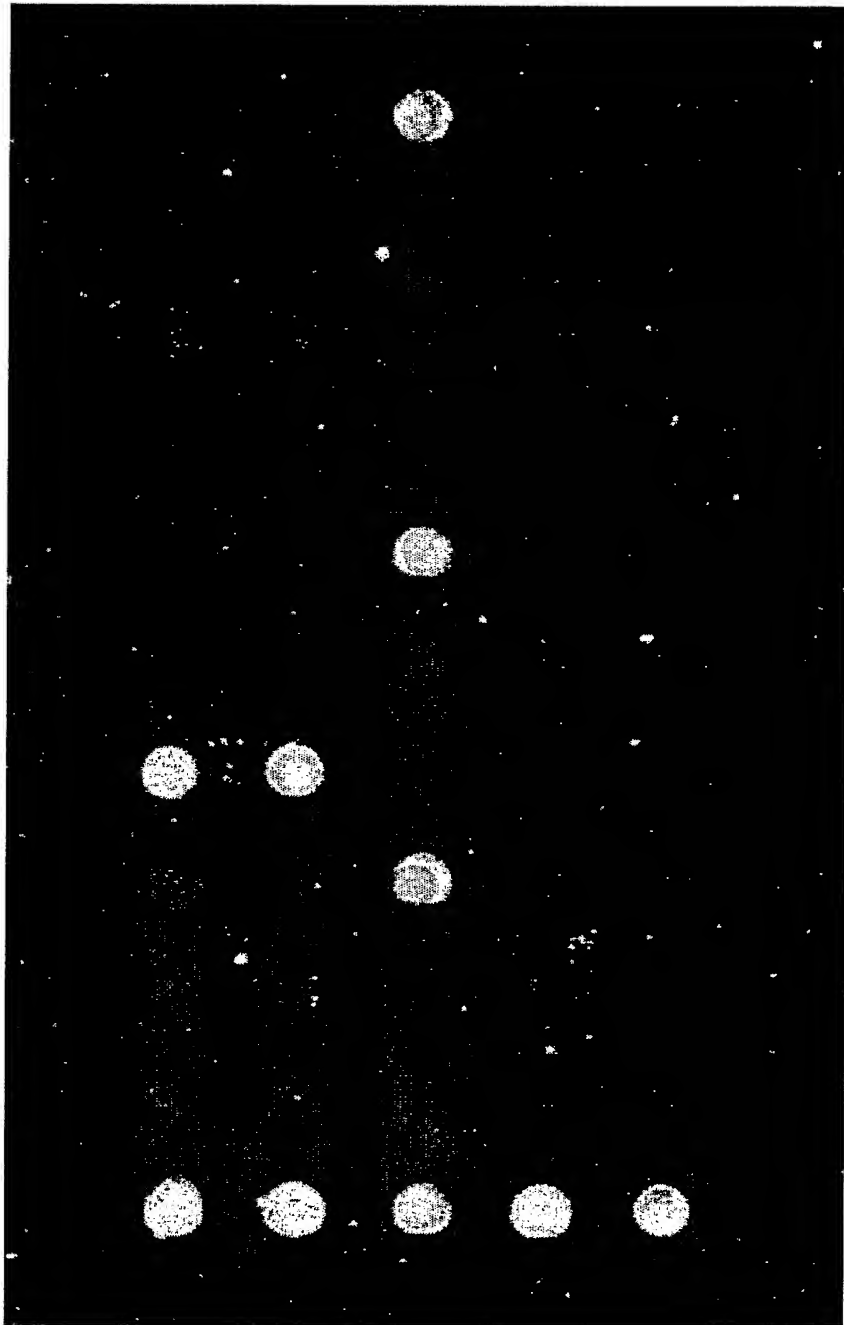


FIG.4

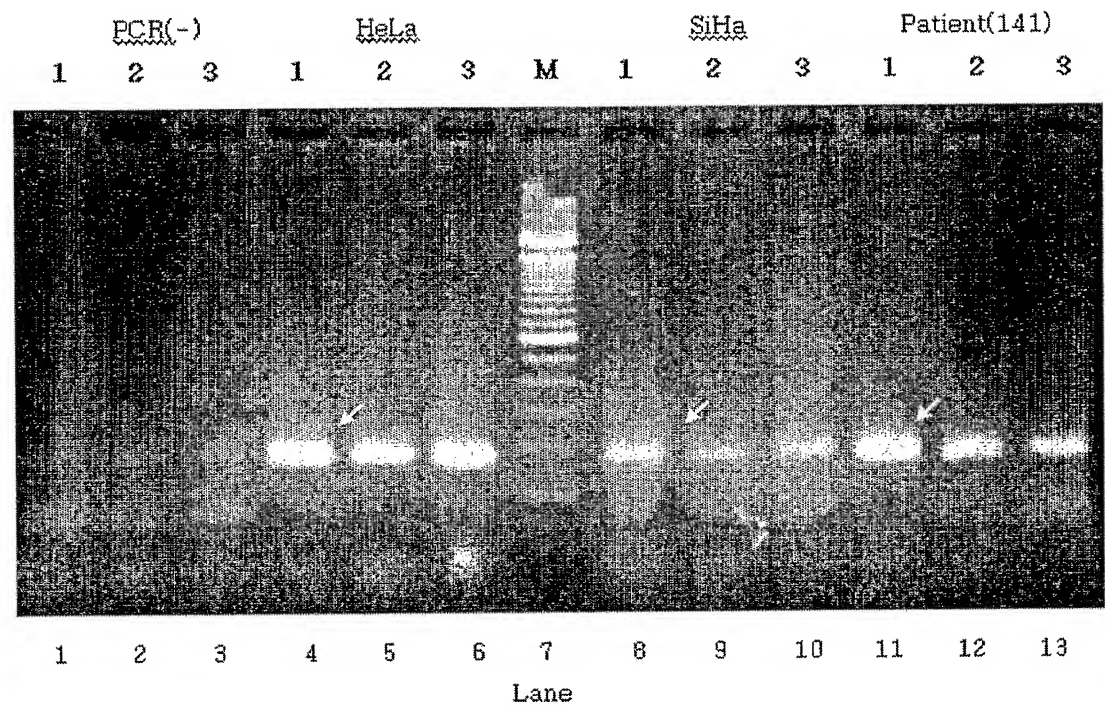


FIG.5

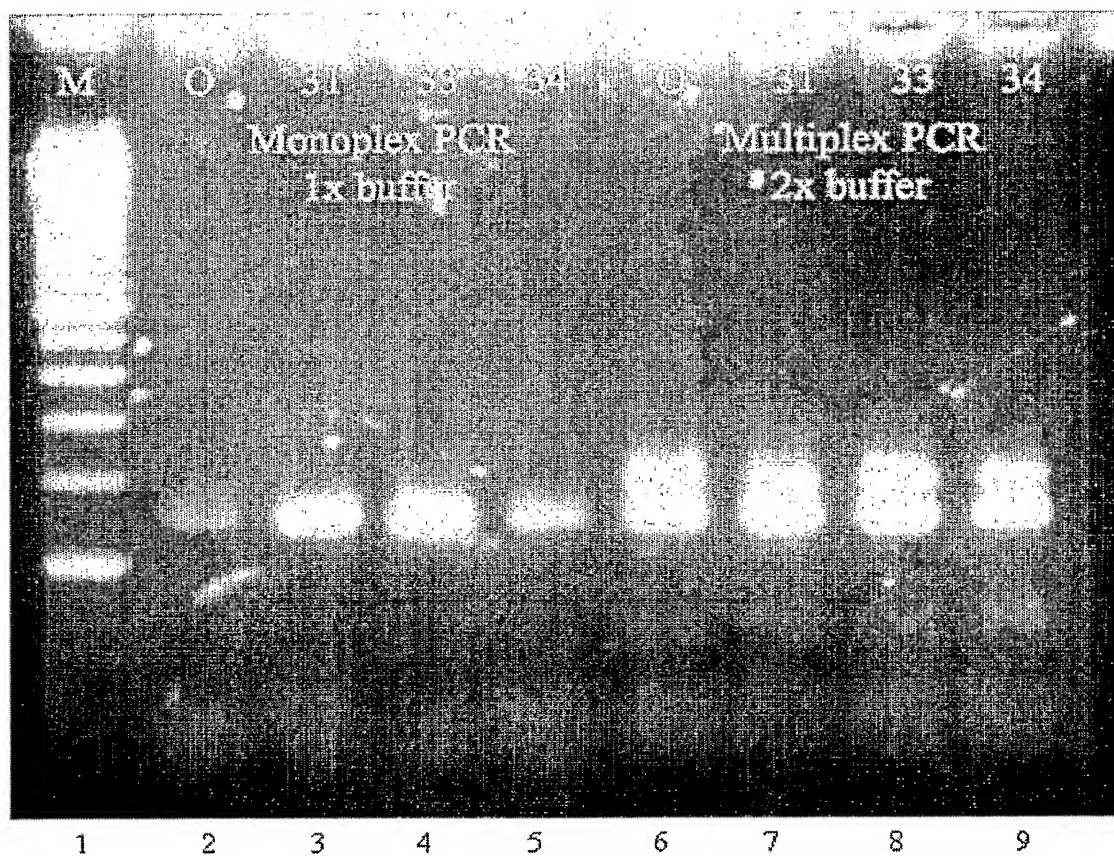


FIG.6

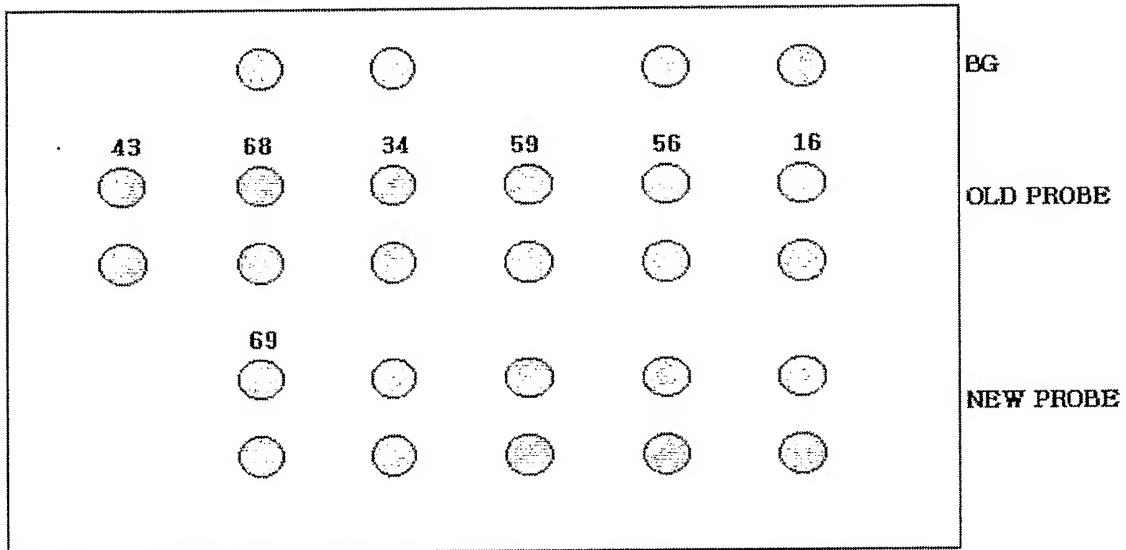


FIG.7A

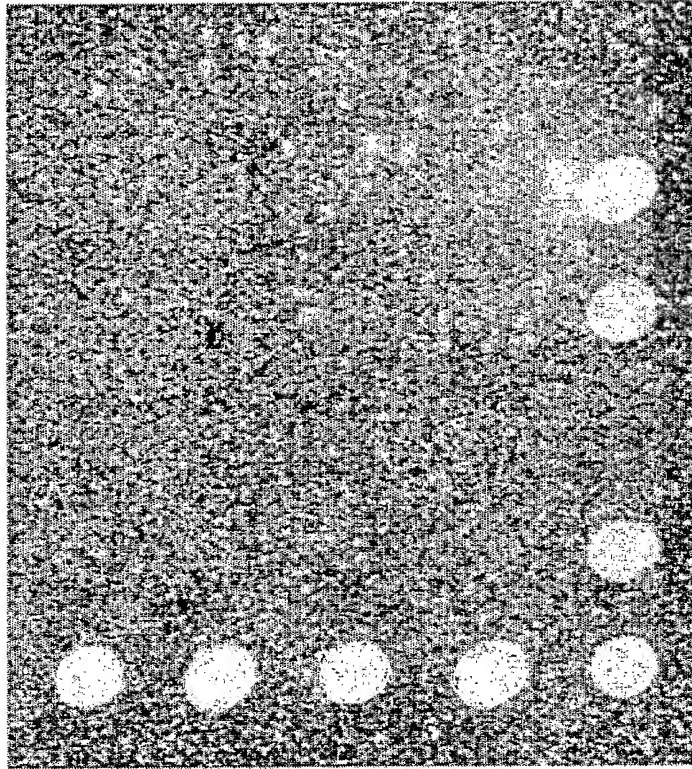


FIG.7B

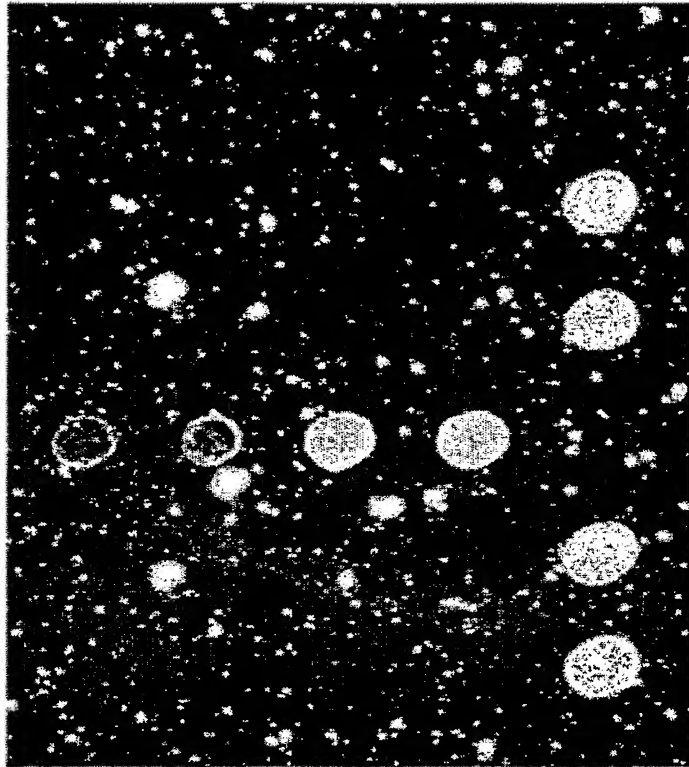


FIG.7C

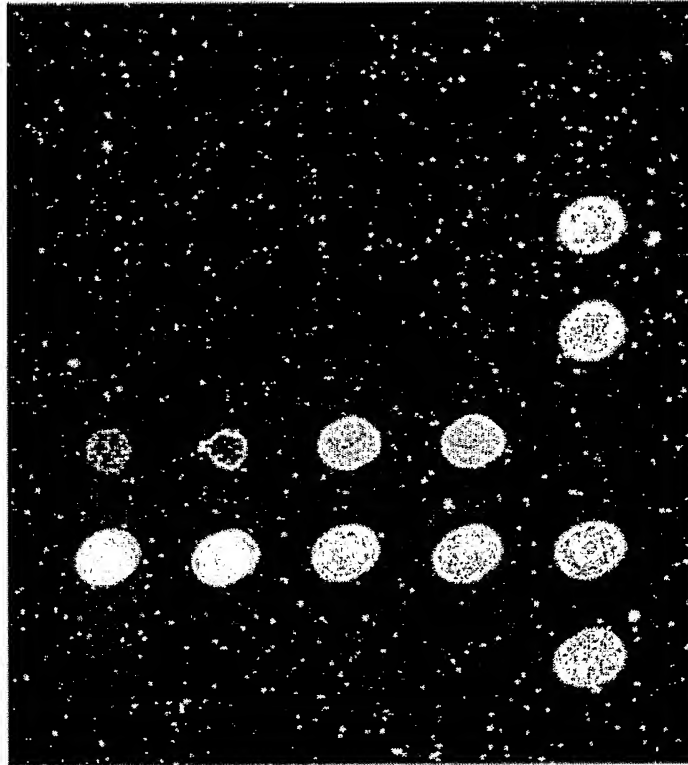


FIG.7D

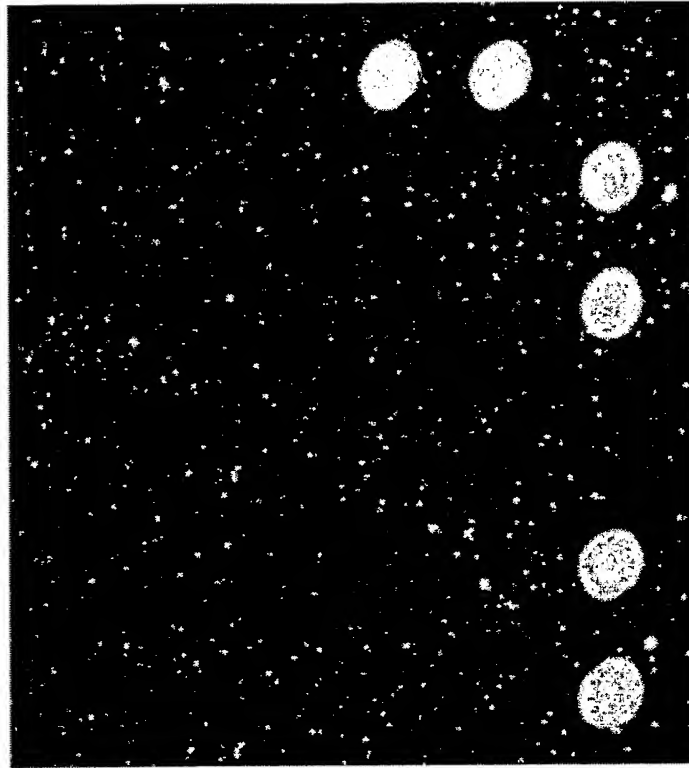


FIG. 7E

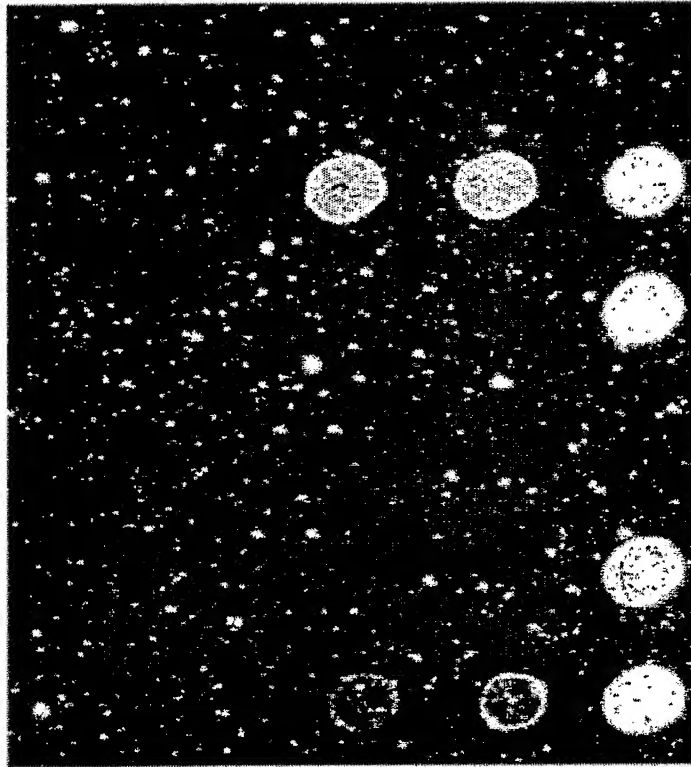
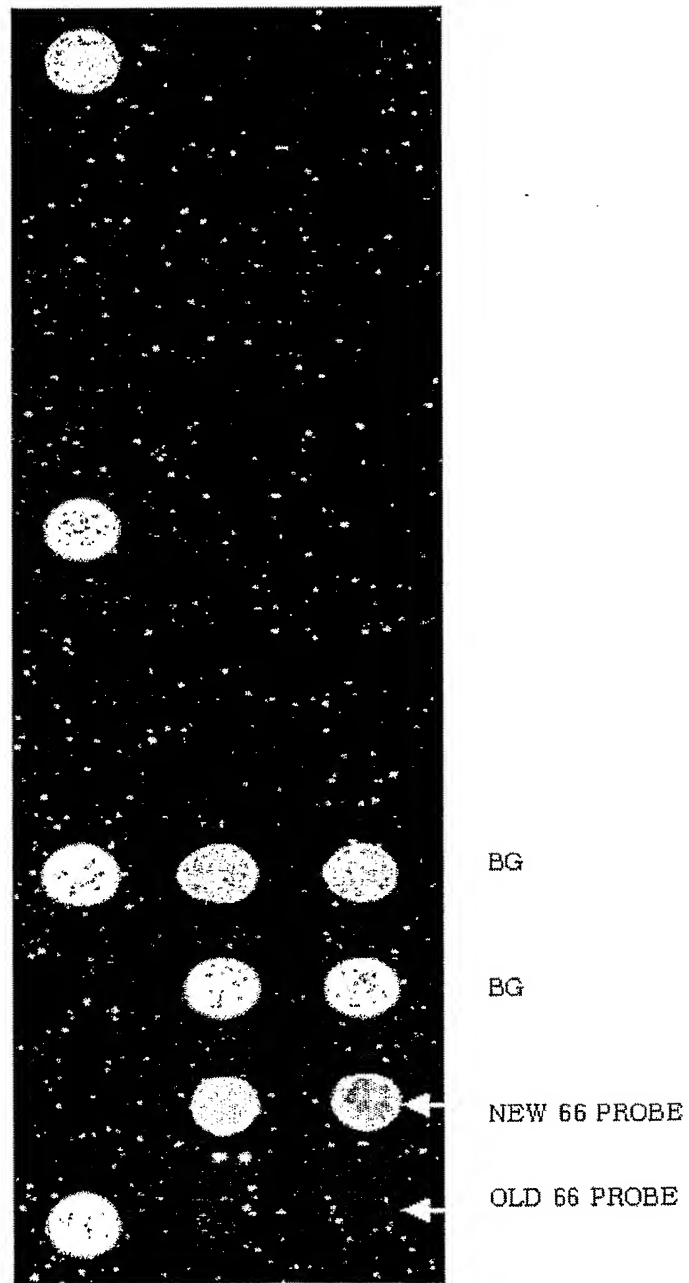


FIG.8



[Sequence Listing]

<110> biomedlab(representative: Kim. Jong-Won)
 <120> Genotyping kit for diagnosis of human papilloma virus infection
 <130> DPP012599
 <160> 54
 <170> Kopatent In 1.71
 <210> 1
 <211> 30
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> porbe HPV 13
 <400> 1
 acatcatctc tttcagacac atataaggcc 30
 <210> 2
 <211> 30
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> probe HPV 26
 <400> 2
 agtacattat ctgcagcatc tgcattccact 30
 <210> 3
 <211> 30
 <212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 30

<400> 3

ttatccacat ataattcaag ccaaattaaa

30

<210> 4

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 43

<400> 4

cctctactga ccctactgtg cccagtacat

30

<210> 5

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> porbe HPV 53

<400> 5

tctacatata attcaaagca aattaaacag ta

32

<210> 6

<211> 31

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 54

<400> 6

tacagcatcc acgcaggata gctttaataa t

31

<210> 7

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 54

<400> 7

catccacgca ggatagcttt aataattctg

30

<210> 8

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 55

<400> 8

tgtgctgcta caactcagtc tccatctaca aca

33

<210> 9

<211> 33

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 57

<400> 9
gaaactaatt ataaagcctc caattataag gaa 33

<210> 10
<211> 31
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 61

<400> 10
ctgtatctga atataagcc acaagcttta g 31

<210> 11
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 61

<400> 11
cctgtatctg aatataaagc cacaagcttt 30

<210> 12
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 62

<400> 12
cctccactgc tgcagcagaa tacacggcta 30

<210> 13
 <211> 33
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> probe HPV 64

<400> 13
 tacaaatcca ccatatgcaa acactaattt taa 33

<210> 14
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> probe HPV 67

<400> 14
 tatgttctga ggaaaaatca gaggtacat 30

<210> 15
 <211> 33
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> probe HPV 68

<400> 15
 ttigtctact actactgaat cagctgtacc aaa 33

<210> 16
 <211> 33

<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 69

<400> 16
aatctgcatc tgccactttt aaaccatcag att 33

<210> 17
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 70

<400> 17
aacggccata cctgctgtat atagccctac 30

<210> 18
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 70

<400> 18
accgaaacgg ccatacctgc tgtatatagc 30

<210> 19
<211> 30
<212> DNA
<213> Artificial Sequence

<220>

<223> probe HPV 74

<400> 19

cgccttctgc tacatataat agttcagact

30

<210> 20

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV JC9710

<400> 20

aaacaccctc tgacacatac aaggcttcca

30

<210> 21

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 16

<400> 21

gtgctgccat atctacttca gaaactacat

30

<210> 22

<211> 34

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 16

<400> 22
tatgtgctgc catatctact tcagaaacta cata 34

<210> 23
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 34

<400> 23
ggtacacaat ccacaagtac aactgcacca 30

<210> 24
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 35

<400> 24
ttctgctgtg tcttctagt acagtacata 30

<210> 25
<211> 36
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 35

<400> 25
gcacggctctg tgtgttctgc tgtgtcttct acgtgc 36

<210> 26
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 40

<400> 26
cttatgtgct gccacacagt cccccacacc 30

<210> 27
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 56

<400> 27
tattagtact gctacagaac agttaagtaa 30

<210> 28
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 58

<400> 28
cactgaagta actaaggaag gtacatataa 30

<210> 29

<211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> probe HPV 59

<400> 29
 tctactactt cttctattcc taatgtatac 30

<210> 30
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> probe HPV 66

<400> 30
 ctaaaagcac attaactaaa tatgatgccc 30

<210> 31
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> probe HPV 6

<400> 31
 atccgtaact acatcttcca catacaccaa 30

<210> 32
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>

<223> probe HPV 11

<400> 32

atctgtgtct aaatctgcta catacactaa

30

<210> 33

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 18

<400> 33

tgcttctaca cagtctcctg tacctgggca

30

<210> 34

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 31

<400> 34

tgttttgtgct gcaattgcaa acagtgatac

30

<210> 35

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 33

<400> 35
tttatgcaca caagtaacta gtgacagtac 30

<210> 36
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 39

<400> 36
tctacctcta tagagtcttc cataccttct 30

<210> 37
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 42

<400> 37
ctgcaacatc tggtgatata tatacagctg 30

<210> 38
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 44

<400> 38

gccactacac agtcccctcc gtctacatat

<210> 39

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 45

<400> 39

acacaaaatc ctgtgccaag tacatatgac 30

<210> 40

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 51

<400> 40

agcactgcca ctgctgcggt ttccccaaca 30

<210> 41

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 52

<400> 41

tgctgagggtt aaaaaggaaa gcacatatataa 30

<210> 42
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> b-globin

<400> 42
tgcacctgac tcctgaggag aagtctgccg 30

<210> 43
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> primer Gp5+

<400> 43
tttkttachg tkgtdgatac yac 23

<210> 44
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> primer Gp6+

<400> 44
gaaahataaa ytgyaadtc taylorc 25

<210> 45
<211> 18
<212> DNA

<213> Artificial Sequence

<220>

<223> primer BG 1

<400> 45

atacaagtca gggcagag

18

<210> 46

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> primer BG 2

<400> 46

cttaaacctg tcttgtaacc

20

<210> 47

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 16

<400> 47

gtcattatgt gctgccatat ctacttcaga

30

<210> 48

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 34

<400> 48

tacacaatcc acaagtacaa atgcaccata

30

<210> 49

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 35

<400> 49

gtctgtgtgt tctgctgtgt cttctagtga

30

<210> 50

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 40

<400> 50

gctgccacac agtccccac accaacccca

30

<210> 51

<211> 30

<212> DNA

<213> Artificial Sequence

<220>

<223> probe HPV 56

<400> 51
gtactgctac agaacagtta agtaaataatg 30

<210> 52
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 58

<400> 52
attatgcact gaagtaacta aggaaggtac 30

<210> 53
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 59

<400> 53
ctgtgtgtgc ttctactact gcttctattc 30

<210> 54
<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> probe HPV 66

<400> 54
ctattaatgc agctaaaagc acattaacta 30

INTERNATIONAL SEARCH REPORT

International application No.

PCT/KR01/01562

A. CLASSIFICATION OF SUBJECT MATTER**IPC7 C12Q 1/68**

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC7 C12Q 1/68

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Korean Patents and applications for inventions since 1975

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

CAPLUS, MEDLINE, BIOSIS, WPINDEX

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X	WO 95/22626 A1 (STICHTING RESEARCHFONDS PATHOLOGIE) 24 AUGUST 1995 (see page 17)	1
A	US 5484699 A (ABOTT LABORATORIES) 16 JANUARY 1996	1 - 14
A	US 5861242 A (AFFYMETRIX, INC) 19 JANUARY 1999	1 - 14

☐ Further documents are listed in the continuation of Box C.☒ See patent family annex.

* Special categories of cited documents:

"A" document defining the general state of the art which is not considered to be of particular relevance

"E" earlier application or patent but published on or after the international filing date

"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of citation or other special reason (as specified)

"O" document referring to an oral disclosure, use, exhibition or other means

"P" document published prior to the international filing date but later than the priority date claimed

"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art

"&" document member of the same patent family


Date of the actual completion of the international search

20 JUNE 2002 (20.06.2002)

Date of mailing of the international search report

22 JUNE 2002 (22.06.2002)

Name and mailing address of the ISA/KR



Korean Intellectual Property Office
920 Dunsan-dong, Seo-gu, Daejeon 302-701,
Republic of Korea

Facsimile No. 82-42-472-7140

Authorized officer

HAN, Hyun Sook

Telephone No. 82-42-481-5596



INTERNATIONAL SEARCH REPORT

Information on patent family members

International application No.

PCT/KR01/01562

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO 95/22626 A1	24.08.95	CA 2183758 AA	24.08.95
		AU 685233 B2	15.01.98
		JP 9509062 T2	16.09.97
US 5484699 A	16.01.96	JP 4281791 A2	07.10.92
		EP 477972 A3	20.05.92
US5861242 A	19.01.99	EP 7300663 A4	23.02.00
		JP 9507121 T2	22.07.97
		AU 8126694	22.05.95